

SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT: Quint, Wilhelmus
Van Doorn, Leendert

(ii) TITLE OF INVENTION: Probes, methods and kits for detection and
typing of Helicobacter pylori nucleic acids in biological
samples.

(iii) NUMBER OF SEQUENCES: 280

(iv) CORRESPONDENCE ADDRESS:

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(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.30

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(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: EP96/870131.8
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(A) APPLICATION NUMBER: PCT/EP97/05614

(B) FILING DATE: 10-OCT-1997

(2) INFORMATION FOR SEQ ID NO: 1:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

GTTGATAACG CTGTCGCTTC

20

(2) INFORMATION FOR SEQ ID NO: 2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

GGAGCRTTRG TCAGCATCAC

20

(2) INFORMATION FOR SEQ ID NO: 3:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 21 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

GCTTTAGTAG GAGCRTTRGT C

21

(2) INFORMATION FOR SEQ ID NO: 4:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

GGAGCGTTGA TTAGYKCCAT

20

(2) INFORMATION FOR SEQ ID NO: 5:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 19 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

GTTTTAGCAG GAGCGTTGA

19

(2) INFORMATION FOR SEQ ID NO: 6:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:

GCTAAYACGC CAAAYGATCC

20

(2) INFORMATION FOR SEQ ID NO: 7:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 20 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

GATCCCATAC ACAGCGAGAG

20

(2) INFORMATION FOR SEQ ID NO: 8:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 19 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

TTGATACGGG TAATGGTGG

19

(2) INFORMATION FOR SEQ ID NO: 9:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 20 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:

GGGTAATGGT GGTTCACA

20

(2) INFORMATION FOR SEQ ID NO: 10:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 21 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:

ACGAATTAA GAGTGAATGG C

21

(2) INFORMATION FOR SEQ ID NO: 11:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 21 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:

AGAGCGATAA CGGGCTAAAC A

21

(2) INFORMATION FOR SEQ ID NO: 12:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 24 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:

TTGACCAACA ACCACAAACC GAAG

24

(2) INFORMATION FOR SEQ ID NO: 13:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 22 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:

CTTCCCTTAA TTGCGAGATT CC

22

(2) INFORMATION FOR SEQ ID NO: 14:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:

CCTGARACCG TTCCTACAGC

20

(2) INFORMATION FOR SEQ ID NO: 15:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:

GTGGATGCYC ATACRGCTWA

20

(2) INFORMATION FOR SEQ ID NO: 16:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:

RTGAGCTTGT TGATATTGAC

20

(2) INFORMATION FOR SEQ ID NO: 17:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 21 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:

CACAGCCACT TTCAATAACG A

21

(2) INFORMATION FOR SEQ ID NO: 18:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 20 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:

CGTCAAAATA ATTCCAAGGG

20

(2) INFORMATION FOR SEQ ID NO: 19:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 19 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:

CAACAACCAC AAACCGAAG

19

(2) INFORMATION FOR SEQ ID NO: 20:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 21 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:

GATTGGTTTT TGATCAGGAT C

21

(2) INFORMATION FOR SEQ ID NO: 21:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21:

GATAAGAAYG ATAGGATAA

20

(2) INFORMATION FOR SEQ ID NO: 22:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 18 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 22:

AATACTGATT CTTTTTGG

18

(2) INFORMATION FOR SEQ ID NO: 23:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 23:

GTGGATGCCC ATACGGCTAA

20

(2) INFORMATION FOR SEQ ID NO: 24:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 24:

GTGGATGCTC ATACAGCTWA

20

(2) INFORMATION FOR SEQ ID NO: 25:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 25:

GTGGATGCCC ATACGATCAA

20

(2) INFORMATION FOR SEQ ID NO: 26:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 20 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 26:

GCGAGCGCTC ATACGGTCAA

20

(2) INFORMATION FOR SEQ ID NO: 27:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 20 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 27:

GGATTTTGA TCGCTTTATT

20

(2) INFORMATION FOR SEQ ID NO: 28:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 20 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 28:

GGGYTATTGG TYAGCATCAC

20

(2) INFORMATION FOR SEQ ID NO: 29:

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 20 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 29:

GCTTTAGTRG GGYTATTGGT

20

(2) INFORMATION FOR SEQ ID NO: 30:

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 19 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 30:

TTGATACKGG TAATGGTGG

19

(2) INFORMATION FOR SEQ ID NO: 31:

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 20 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 31:

KG GTAATGGT GGT TCAACA

20

(2) INFORMATION FOR SEQ ID NO: 32:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 32:

KG GTAATGGT GGT TCAACA

20

(2) INFORMATION FOR SEQ ID NO: 33:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 21 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 33:

AGAGCGATAA YGKCTAAAC A

21

(2) INFORMATION FOR SEQ ID NO: 34:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 21 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 34:

AGGGTAGAAA TGGTATCGAC A

21

(2) INFORMATION FOR SEQ ID NO: 35:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 23 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 35:

CATGCGCCT TCTTTACAAC CGT

23

(2) INFORMATION FOR SEQ ID NO: 36:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 23 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 36:

CATGCCGCCT TTTTCACRAC CGT

23

(2) INFORMATION FOR SEQ ID NO: 37:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 23 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 37:

CATGCCGCTC' TTTTACAAC CGT

23

(2) INFORMATION FOR SEQ ID NO: 38:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 23 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 38:

CATGCCGCCT TTTTACAAC CGT

23

(2) INFORMATION FOR SEQ ID NO: 39:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 23 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 39:

AGTCGCGCYT TTTYACAAC CGT

23

(2) INFORMATION FOR SEQ ID NO: 40:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 184 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 40:

CCCTCTGGTT TCTCTCGCTT TAGTAGGAGC ATTGGTCAGC ATCACACCGC AACAAAGTCA	60
TGCCGCCTTT TTCACAACCG TGATCATTC AGCCATTGTT GGGGGTATCG CTACAGGCAC	120
CGCTGTAGGA ACGGTCTCAG GGCTTCTTAG TTGGGGACTA AAACAAGCCG AAGAAGCCAA	180
TAAA	184

(2) INFORMATION FOR SEQ ID NO: 41:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 199 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 41:

TCGCCCTCTN GTTCTCTCG CTTTAGTAGN AGCATTNGTC AGCATCACAC CGCAACANAG 60

TCATGCCGCC TTTTTCACAA CCGTNATCAT TCCAGCCATT GTTGGGGGTA TNGCTACAGG	120
CACCGCTGTA GGAACGGTCT CAGGGCTTCT TAGTTGGGGA CTAAAACAAG CCGAAGAAGC	180
CAATAAAACC CCAGATAAA	199

(2) INFORMATION FOR SEQ ID NO: 42:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 227 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 42:

GAATCGCCCT YTGGTTTCTC TTGCTTTAGT AGGAGCATTG RTTAGYRYCA YACCGCAACA	60
AAGTCATGCC GCCTTTTYYA CRACCGTGAT CATTCCAGCC ATTGTTGGRG GTATCGCTAC	120
AGGCACTGCT GTAGGAACGG TCTCAGGGCT TCTTAGTTGG GGRCTCAAAC AAGCCGAAGA	180
AGCSAATAAA ACCCCRGATA AACCCGATAA AGTTTGGCGC ATTCAAG	227

(2) INFORMATION FOR SEQ ID NO: 43:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 176 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 43:

GCCCTTTAGT TTCTCTCGCT TTAGTGGGGT TATTGGTCAG CATCACACCA CAAAAAAGTC	60
ATGCCGCCTT TTTCACAACC GTGATCATTC CAGCCATTGT TGGAGGTATC GCTACAGGTG	120
CTGCTGTAGG AACGGTCTCA GGGCTTCTTG GTTGGGGGCT CAAACAAGCC GAAGAA	176

(2) INFORMATION FOR SEQ ID NO: 44:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 185 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 44:

GCCCTCTGGT TTCTCTCGCT TTAGTAGGAG CATTGGTCAG CATCACACCG CAACAAAGTC	60
ATGCCGCCTT TTTCACAACC GTGATCATTC CAGCCATTGT TGGAGGTATC GCTACAGGCG	120
CTGCTGTAGG AACGGTCTCA GGGCTTCTTA GCTGGGGGCT CAAACAAGCC GAAGAAGCCA	180
ATAAA	185

(2) INFORMATION FOR SEQ ID NO: 45:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 204 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 45:

AATCGCCCTC TGGTTTCTCT CGCTTTAGTA GGAGCATTGG TCAGCATCAC ACCGCAACAA	60
AGTCATGCCG CCTTTTTTAC AACCGTGATC ATTCCAGCCA TTGTTGGAGG TATCGCTACA	120
GGCGCTGCTG TAGGAACGGT CTCAGGGCTT CTTAGCTGGG GGCTCAAACA AGCCGAACAA	180
GCCAATAAAG CCCC GGACAA ACCC	204

(2) INFORMATION FOR SEQ ID NO: 46:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 207 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 46:

GAATCGCCCT TTAGTTTCTC TTGCTTTAGT AGGAGCATTG GTCAGCATCA CACCGCAACA	60
AAGTCATGCC GCCTTTTTCA CAACCGTGAT CATTCCAGCC ATTGTTGGGG GTATCGCTAC	120
AGGCGCTGCT GTAGGAACGG TTTCAGGGCT TCTTGGCTGG GGGCTAAAAC AAGCCGAAGA	180
AGCCAATAAA ACCCCAGATA AACCCGA	207

(2) INFORMATION FOR SEQ ID NO: 47:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 207 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 47:

CAATCGCCCT CTAGTTTCTC TCGCTTTAGT AGGAGCATTG GTCAGCATCA CACCGCAACA	60
AAGTCATGCC GCCTTTTTCA CAACCGTGAT CATTCCAGCC ATTGTGGGGG GTATCGCTAC	120
AGGCGCTGCT GTAGGAACGG TCTCAGGGCT TCTTAGCTGG GGGCTCAAAC AAGCCGAAGA	180
AGCCAATAAA ACCCCGGACA AACCCGA	207

(2) INFORMATION FOR SEQ ID NO: 48:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 207 base pairs
 - (B) TYPE: nucleic acid

(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 48:

CAATCGCCCT CTGGTTTCTC TTGCTTTAGT AGGAGCGTTA GTCAGCATCA CACCGCAACA	60
AAGTCATGCC GCCTTTTTCA CAACCGTGAT CATTCCAGCC ATTGTTGGGG GGATCGCTAC	120
AGGCGCTGCT GTAGGAACGG TCTCAGGGCT TCTTAGCTGG GGGCTCAAAC AAGCCGAAGA	180
AGCCAATAAA ACCCCAGATA AACCCGA	207

(2) INFORMATION FOR SEQ ID NO: 49:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 176 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 49:

CCCTTTAGTT TCTCTTGTTT TAGCAGGAGC GTTGATTAGC GCCATACCGC AACAAAGTCA	60
TGCCGCCCTTT TTCACGACCG TGATCATTCC AGCCATTGTT GGGGGTATCG CTACAGGCAC	120
CGCTGTAGGA ACGGTTTCAG GGCTTCTTAG CTGGGGGCTC AAACAAGCCG AAGAAG	176

(2) INFORMATION FOR SEQ ID NO: 50:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 187 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 50:

CCCTTTAGTT TCTCCTGTTT TAGCAGGAGC GTTGATTAGC TCCATACCGC AACAAAGTCA	60
TGCCGCCTTT TTCACAACCG TGATCATTC AGCCATTGTT GGGGGTATCG CTACAGGCAC	120
TGCTGTAGGA ACGGTCTCAG GGCTTCTTAG CTGGGGGCTC AAACAAGCYG AASAAGCSAA	180
TAAAGCC	187

(2) INFORMATION FOR SEQ ID NO: 51:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 193 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 51:

CTTGAGTTTC TCTTGTTTTA GCAGGAGCGT TGATTAGCGC CATACCGCAA CAAAGTCATG	60
CCGCCTTTTT CACGACCGTG ATCATTCCAG CCATTGTTGG GGGTATCGCT ACAGGCACCG	120
CTGTAGGAAC GGTTCAGGG CTTCTTAGCT GGGGGCTCAA ACAAGCCGAA GAAGCCAATA	180
AAACCCCA TAA	193

(2) INFORMATION FOR SEQ ID NO: 52:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 196 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 52:

GCCCTTTAGT TTCTCTCGTT TTAGCAGGAG CGTTGATTAG CTCCATACCG CAACAAAGTC	60
ATGCCGCCTT TTTCACAACC GTGATCATTC CAGCCATTGT TGGGGGTATC GCTACAGGCA	120
CCGCTGTAGG AACGGTTTCA GGGCTTCTTA GCTGGGGGCT CAAACAAGCC GAACAAGCCA	180
ATAAAGCCCC GGACAA	196

(2) INFORMATION FOR SEQ ID NO: 53:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 131 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 53:

AATCGCCCTT TAGTTTCTCT TGTTTTCAGCA GGAGCGTTGA TTAGCGCCAT ACCGCAACAA	60
AGTCATGCCG CCTTTTTCAC GACCGTNATC ATTCCAGCCA TTGTTNNNNG TATCGCTACA	120
GGCACCGCTG T	131

(2) INFORMATION FOR SEQ ID NO: 54:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 185 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 54:

CTACGCCCTT TAGTTTCTCT CGTTTTAGCA GGAGCGTTGA TTAGCGCCAT ACCGCAACAA	60
AGTCATGCCG CCTTTTTTCAC AACCGTGATC ATTCCAGCCA TTGTTGGGGG CATCGCTTCA	120
GGCGCTGCTG TAGGAACGGT CTCAGGGCTT CTTAGTTGGG GACTCAAACA AGCCGAAGAA	180
GCGAA	185

(2) INFORMATION FOR SEQ ID NO: 55:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 201 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 55:

AATCGCCCTT TAGTTTCTCT CGTTTTAGCA GGAGCGTTGA TTAGCGCCAT ACCGCAAGAG	60
AGTCATGCCG CCTTTTTTCAC AACCGTNATC ATTCCAGCCA TTGTTGGGGG TATCGCTACA	120
GGCACCCTG TAGGAACGGT CTNAGGGCTT YTTAGTTGGG GACTNWAACA AGCCGAAGAA	180
GCCAATAAAA CCCCCGATAA A	201

(2) INFORMATION FOR SEQ ID NO: 56:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 187 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 56:

CCCTTTAGTT TCTCTCGTTT TAGCANNAGC GTTGATTAGC RCCATACCGC AAGAGAGTCA	60
---	----

TGCCGCCTTT TTTACAACCG TNATCATTC AGCCATTGTT GGGGGTATCG CTACAGGYGC	120
TGCTGTAGGA ACGGTCTCAG GGCTTCTTAG CTGGGGGCTC AAACAAGCCG AACAAGCCAA	180
TAAAGCC	187

(2) INFORMATION FOR SEQ ID NO: 57:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 187 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 57:

CCCTTTGGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC TCCATACCGC AAGAGAGTCA	60
TGCCGCCTTT TTCACAACCG TGATCATTC AGCCATTGTT GGGGGTATCG CTACAGGCGC	120
TGCTGTAGGA ACGGTCTCAG GGCTTCTTAG CTGGGGGCTC AAACAAGCCG AAGAAGCGAA	180
TAAAACC	187

(2) INFORMATION FOR SEQ ID NO: 58:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 187 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 58:

CCCTTTAGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC TCCATACCGC AAGAGAGTCA	60
TGCCGCCTTT TTCACAACCG TGATCATTC AGCCATTGTT GGGGGTATCG CTACAGGCAC	120
CGCTGTAGGA ACGGTCTCAG GGCTTCTTAG CTGGGGACTC AAACAAGCCG AAGAAGCGAA	180

TAAAACC

187

(2) INFORMATION FOR SEQ ID NO: 59:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 185 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 59:

CCCTTTAGTT TCTCTCGTTT TAGCAGNAGC GTTNATTAGT GCCATACCGC AANANAGTCA	60
TGCCGCCTTT TTCACNACCG TNATCATTC AGCCATTGTT GGGGGTATCG CCACAGGCAC	120
CGCTGTAGGA ACGGTCTCAG GGCTTCTTAG TTGGGGACTN AAACAAGCCG AAGAAGCGAA	180
TAAAA	185

(2) INFORMATION FOR SEQ ID NO: 60:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 199 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 60:

CAGGAGCGTT GATTAGTGCC ATACCGCAAG AGAGTCATGC CGCCTTTTTC ACGACCGTGA	60
TCATTCCAGC CATTGTTGGG GGTATCGCCA CAGGCACCGC TG TAGGAACG GTCTCAGGGC	120
TTCTTAGTTG GGGACTCAAA CAAGCCGAAG AAGCGAATAA AACCCAGTA TAAACCCGAT	180
AAAGTTTGGC GCATTCAAG	199

(2) INFORMATION FOR SEQ ID NO: 61:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 188 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 61:

TTTCTCTCGT TTTAGCAGGA GCGTTGATTA GCTCCATACC GCAAGAGAGT CATGCCGCCT	60
TTTTCACAAC CGTGATCATT CCAGCCATTG TTGGGGGTAT CGCTACAGGC GCTGCTGTAG	120
GAACGGTCTC AGGGCTTCTT AGCTGGGGGC TCAAACAAGC CGAAGAAGCG AATAAAACCC	180
CAGATAAA	188

(2) INFORMATION FOR SEQ ID NO: 62:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 206 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 62:

GAATCGCCCT TTAGTTTCTC TCGTTTTAGC AGGAGCGTTG ATTAGTGCCA TACCGCAAGA	60
GAGTCATGCC GCCTTTTCA CGACCGTGAT CATTCCAGCC ATTGTTGGGG GTATCGCCAC	120
AGGCACCGCT GTAGGAACGG TCTCAGGGCT TCTTAGTTGG GGA CTCAAAC AAGCCGAAGA	180
AGCGAATAAA ACCCCAGTAT AAACCC	206

(2) INFORMATION FOR SEQ ID NO: 63:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 206 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 63:

AATCGCCCTT TGGTTTCTCT CGTTTTCAGCA GGAGCGTTGA TTAGCTCCAT ACCGCAAGAG	60
AGTCATGCCG CCTTTTTCAC AACCGTGMTC ATTCCAGCCA TTGTTGGGGG TATGGCTACA	120
GGCGCTGCTG TAGGAACGGT CTNAGGGCTT YTTAGCTGGG GGCTNWAACA AGCCGAAGAA	180
GCGAATAAAA CCCAGATAA ACCCGA	206

(2) INFORMATION FOR SEQ ID NO: 64:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 201 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 64:

AATCGCCCTA TTATTTCTCT CGTTTTCAGCA GGAGCGTTGA TTAGCTCCAT ACCGCAAGAA	60
AGTCATGCCG CCTTTTTCAC GACCGTGATC ATTCCAGCCA TTGTTGGGGG TATCGCCACA	120
GGCGCTGCTG TAGGAACGGT CTCAGGGCTT CTTAGCTGGG GGCTCAAACA AGCCGAACAA	180
GCCAATAAAG CCCCGGACAA A	201

(2) INFORMATION FOR SEQ ID NO: 65:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 197 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 65:

```
GCCCTTTAGT TTCTCTTGTT TTAGCAGGAG CGTTGATTAG TGCCATACCG CAAGAGAGCT      60
ATGCCGCCTT TTTCACAACC GTGATCATTC CAGCCATTGT TGGGGGTATC GCTACAGGCA      120
CCGCTGTAGG AACGGTCTCA GGGCTTCTTA GTTGGGGACT CAAACAAGCC GAAGAAGCGA      180
ATAAAACCCC AGATAAA                                          197
```

(2) INFORMATION FOR SEQ ID NO: 66:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 201 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 66:

```
AATCGCCCTT TAGTTTCTCT CGTTTTAGCA GGAGCGTTTA TTAGCGCCAT ACCGCAAGAG      60
AGTCATGCCG CCTTTTTCAC GACCGTGATC ATTCCAGCCA TTGTTGGGGG TATCGCCACA      120
GGCACCGCTG TAGGAACGGT CTCAGGGCTT CTTAGTTGGG GACTCAAACA AGCCGAAGAA      180
GCGAATAAAA CCCCAGATAA A                                          201
```

(2) INFORMATION FOR SEQ ID NO: 67:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 195 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 67:

GTGGGGGTGT TAATGGGCAC CGAACTAGGG GCTAATACGC CAAACGATCC CATAACAGC	60
GAGAGTCGCG CCTTTTTCAC AACCGTGATC ATTCCAGCCA TTGTTGGGGG TATCGCCACA	120
GGCACTGCTG TAGGAACGGT CTCAGGGCTT CTTAGTTGGG GACTCAAACA AGCCGAAGAA	180
GCGAATAAAAA CCCCCA	195

(2) INFORMATION FOR SEQ ID NO: 68:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 196 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 68:

AGTGGGGGTG TTAATGGGCA CTGAACTAGG GGCTAACACG CCAAACGATC CCATAACAG	60
CGAGAGTCGC GCCTTTTTTA CAACCGTGAT CATTCCAGCC ATTGTTGGGG GTATCGCTAC	120
AGGCGCTGCT GTAGGAACGG TTTCAGGGCT TCTTAGCTGG GGGCTCAAAC AAGCCGAACA	180
AGCCAATAAAA GCCCCG	196

(2) INFORMATION FOR SEQ ID NO: 69:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 196 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 69:

TAGTGGGGGC ATTAATGAGT ACCGAACTAG GGGCTAACAC GCCAAATGAT CCCATACACA	60
GCGAGAGTCG CGCCTTTTTT ACAACCGTGA TCATTCCAGC CATTGTTGGG GGTATCGCTA	120
CAGGCGCTGC TGTAGGAACG GTCTCAGGGC TTCTTAGTCG GGGGCTCAAA CAAGCCGAAC	180
AAGCCAATAA AGCCCC	196

(2) INFORMATION FOR SEQ ID NO: 70:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 232 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 70:

CAATCGCCCT ATTATCTCTC TCGCTCTAGT GGGGGTGTTA ATGGGTACCG AACTAGGGGC	60
TAACACGCCA AACGATCCCA TACACAGCGA GAGTCGCGCC TTTTTTACAA CCGTGATCAT	120
TCCAGCCATT GTTGGGGGTA TCGCTACAGG CGCTGCTGTA GGAACGGTTT CAGGGCTTCT	180
TAGCTGGGGG CTCAAACAAG CCGAACAAGC CAATAAAGCC CCGGACAAAC CC	232

(2) INFORMATION FOR SEQ ID NO: 71:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 228 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 71:

AATCGCCCTA TTATCTCTCT CGCTCTAGTG GGGGTGTTAA TGGGTACCGA ACTAGGGGCT	60
AACACGCCAA ACGATCCCAT ACACAGCGAG AGTCGCGCCT TTTTCACAAC CGTGATCATT	120
CCAGCCATTG TTGGAGGTAT CGCTACAGGT GCTGCTGTAG GAACGGTCTC AGGGCTTCTT	180
AGCTGGGGGC TCAAACAAGC CGAACAAGCC AATAAAGCCC CGGACAAA	228

(2) INFORMATION FOR SEQ ID NO: 72:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 228 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 72:

AATCGCCCTA TTATCTCTCT CGCTTTAGTG GGGGTRTTAA TGGGCACCGA ACTAGGGGCT	60
AACACGCCAA ACGATCCCAT ACACAGCGAG AGTCGCGCCT TTTTCACAAC CGTGATCATT	120
CCAGCCATTG TTGGGGGTAT CGCTACAGGC GCTGCTGTAG GAACGGTCTC AGGGCTTCTT	180
AGCTGGGGGC TCAAACAAGC CGAACAAGCC AATAAAGCCC CGGATAAA	228

(2) INFORMATION FOR SEQ ID NO: 73:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 233 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 73:

AATCGCCCTA TTATCTCTCT CGCTTTAGAG GGGGTGTAA TAGGCACCGA ACTAGGGGCT	60
AACACGCCAA ATGATCCCAT ACACAGCGAG AGTCGCGCCT TTTTACAAC CGTTATTATT	120
CCAGCCATTG TTGGGGGTAT CGCTACAGGC GCTGCTGTAG GAACGGTCTC AGGGCTTCTT	180
AGCTGGGGGC TCAAACAAGC CGAACAAGCC AATAAAGCCC CGGATAAACC CGA	233

(2) INFORMATION FOR SEQ ID NO: 74:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 300 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 74:

TTTAAAGGTG GATGCTCATA CAGCTAATTT TAAAGGTATT GATACGGGTA ATGGTGGTTT	60
CAACACCTTA GATTTTAGTG GTGTTACAGG TAAGGTCAAT ATCAACAAGC TCATCACAGC	120
TTCCACTAAT GTGGCCGTTA AAAACTTCAA CATTAAATGAA TTGATTGTTA AAACCAATGG	180
TGTGAGTGTG GGGGAATACA CTCATTTTAG CGAAGATATA GGCAGTCAAT CGCGCATCAA	240
TACCGTGCCT TTGGAAACTG GCACTAGGTC AATCTTTTCT GGGGGTGTTA AATTAAAGG	300

(2) INFORMATION FOR SEQ ID NO: 75:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 300 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 75:

TTTAAAGGTG GATGCTCATA CAGCTAATTT TAAAGGTATT GATACGGGTA ATGGTGGTTT	60
CAACACCTTA GATTTTAGTG GTGTTACAGG TAAGGTCAAT ATCAACAAGC TCATTACGGC	120
TTCCACTAAT GTGGCCGTTA AAAACTTCAA CATTAAATGAA TTGTTGGTTA AGACCAATGG	180
GGTGAGTGTG GGGGAATACA CTCATTTTAG CGAAGATATA GGCAGTCAAT CGCGCATCAA	240
TACCGTGCCT TTGGAAACTG GCACTAGGTC AATCTTTTCT GGGGGTGTCA AATTTAAAGG	300

(2) INFORMATION FOR SEQ ID NO: 76:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 300 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 76:

TTTAAAGTG GATGCTCATA CAGCTAATTT TAAAGGTATT GATACGGGTA ATGGTGGTTT	60
CAACACCTTG GATTTTAGTG GCGTTACAGA CAAAGTCAAT ATCAACAAGC TCATCACAGC	120
TTCCACTAAT GTGGCCATTA AAAACTTCAA CATTAAATGAA TTGTTGGTTA AGACCAATGG	180
GGTGAGTGTG GGGGAATACA CTCATTTTAG CGAAGATATA GGCAGTCAAT CGCGCATCAA	240
CACCGTGCCT TTAGAAACTG GCACTAGGTC AATCTTTTCT GGGGGTGTCA AATTTAAAAG	300

(2) INFORMATION FOR SEQ ID NO: 77:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 300 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 77:

TTTAAAGGTG GATGCTCATA CAGCTAATTT TAAAGGTATT GATACGGGTA ATGGTGGTTT	60
CAACACCTTA GATTTTAGTG GTGTTACAGG TAAGGTCAAT ATCAACAAGC TCATCACAGC	120
TTCCACTAAT GTGGCCGCTA AAAACTTCAA CATTAATGAA TTGATTGTTA AAACCAATGG	180
GGTGAGTGTG GGGGAATACA CTCATTTTAG CGAAGATATA GGCAGTCAAT CGCGCATCAA	240
TACCGTGCGT TTGGAAACTG GCACTAGGTC AATCTATTCT GGCGGTGTTA AATTAAAGG	300

(2) INFORMATION FOR SEQ ID NO: 78:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 300 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 78:

TTTAAAGGTG GATGCTCATA CAGCTAATTT TAAAGGTATT GATACGGGTA ATGGTGGTTT	60
CAACACCTTA GATTTTAGTG GTGTTACAGG TAAGGTCAAT ATCAACAAGC TCATCACAGC	120
TTCCACTAAT GTGGCCGCTA AAAACTTCAA CATTAATGAA TTGATTGTTA AAACCAATGG	180
GGTGAGTGTG GGGGAATACA CTCATTTTAG CGAAGATATA GGCAGTCAAT CGCGCATCAA	240
TACCGTGCGT TTGGAAACTG GCACTAGGTC AATCTATTCT GGCGGTGTTA AATTAAAGG	300

(2) INFORMATION FOR SEQ ID NO: 79:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 300 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 79:

TTTAAAAGTG GATGCTCATA CAGCTAATTT TAAAGGTATT GATACTGGTA ATGGTGGTTT	60
CAACACCTTA GATTTTAGTG GTGTTACAAA CAAAGTCAAT ATCAACAAGC TCATTACAGC	120
TTCCACTAAT GTGGCCGTTA AAAACTTCAA CATTAAATGAA TTGTTGGTTA AGATTAATGG	180
GGTGAGTGTG GGGGAATACA CTTATTTTAG CGAAGATATA GGCAGTCAAT CGCGCATCAA	240
CACCGTGCGT TTGGAAACTG GCACTAGGTC AATCTATTCT GGCGGTGTTA AATTTAAAGG	300

(2) INFORMATION FOR SEQ ID NO: 80:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 300 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 80:

TTTAAAGGTG GATGCTCATA CAGCTAATTT TAAAGGTATT GATACGGGTA ATGGTGGTTT	60
CAACACCTTA GATTTTAGTG GTGTTACAGG TAAGGTCAAT ATCAACAAGC TCATCACGGC	120
TTCCACTAAT GTGGCCGTTA AAAACAACAA CATTAAATGAA TTGGTGGTTA AAACCAATGG	180
GATAAGTGTG GGGGAATACA CTCATTTTAG CGAAGATATA GGCAGTCAAT CGCGCATCAA	240
TACCGTGCGT TTGGAAACAG GCACTAGGTC AATCTTTTCT GGGGGTGTCA AATTTAAAAG	300

(2) INFORMATION FOR SEQ ID NO: 81:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 300 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 81:

TTTAAAAGTG GATGCTCATA CAGCTAATTT TAAAGGTATT GATACGGGTA ATGGTGGTTT	60
CAACACCTTA GATTTTAGTG GTGTTACAGG TAAGGTCAAT ATCAACAAGC TCATTACGGC	120
TTCCACTAAT GTAGCCGTTA AAAACTTCAA CATTAATGAA TTGTTGGTTA AGACCAATGG	180
GGTGAGTGTG GGGGAATACA CTCATTTTAG CGAAGATATA GGCAGTCAAT CGCGCATCAA	240
CACCGTGCGT TTGGAAACTG GCACTAGGTC AATCTTTTCT GGGGGTGTCA AATTAAAAAG	300

(2) INFORMATION FOR SEQ ID NO: 82:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 300 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 82:

TTTAAAGGTG GATGCTCATA CAGCTAATTT TAAAGGTATT GATACGGGTA ATGGTGGTTT	60
CAACACCTTA GATTTTAGTG GTGTTACAGG TAAGGTCAAT ATCAACAAGC TCATCACAGC	120
TTCCACTAAT GTGGCCGTTA AAAACTTCAA CATTAATGAA TTGATTGTTA AAACCAATGG	180
GATAAGTGTG GGGGAATACA CTCATTTTAG CGAAGATATA GGAAGTCAAT CGCGCATCAA	240
TACCGTGCGT TTGGAAACTG GCACTAGATC AATCTTTTCT GGGGGTGTTA AATTAAAGG	300

(2) INFORMATION FOR SEQ ID NO: 83:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 375 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 83:

TTTAAGAGTG GACGCTCATA CAGCTTATTT TAATGGCAAT ATTTATCTGG GAAAATCCAC	60
GAATTTAAGA GTGAATGGCC ATAGCGCTCA TTTTAAAAAT ATTGATGCCA GCAAGAGCGA	120
TAACGGGCTA AACACTAGCG CTTTGGATTT CAGCGGCGTT ACAGACAAAG TCAATATCAA	180
CAAGCTCACT ACATCTGCCA CTAATGTGAA CGTTAAAAAC TTTGACGTTA AGGAATTGGT	240
GGTTACAACC CGTGTCAGA GTTTTGGGCA ATACACTATT TTTGGCGAAA ATATAGGCGA	300
TAAGTCTCGC ATTGGTGTCTG TGAGTTTGCA AACGGGATAT AGCCCGGCCT ATTCTGGGGG	360
CGTTACTTTT AAAAG	375

(2) INFORMATION FOR SEQ ID NO: 84:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 375 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single .

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 84:

CTTAAGAGTG GATGCTCATA CAGCTTATTT TAATGGCAAT ATTTATTTGG GAAAATCCAC	60
GAATTTAAGA GTGAATGGCC ATAGCGCTCA TTTTAAAAAT ATTGATGCCA GTAAGAGCGA	120
TAACGGGCTA AACACTAGTG CTTTGGATTT TAGCGGCGTT ACAGATAAAG TCAATATCAA	180
CAAGCTCACT ACATCTGCCA CTAATGTGAA CGTTAAAAAC TTTGACATTA AGGAATTGGT	240
GGTTACAACC CGAGTTCAAA GTTTTGGGCA ATACACTATT TTTGGCGAAA ATATAGGCGA	300
TAAGTCTCGC ATTGGTGTCTG TTAGTTTGCA AACGGGATAT AGCCCGGCCT ATTCTGGGGG	360

CGTTACTTTT AAAAG

375

(2) INFORMATION FOR SEQ ID NO: 85:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 374 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 85:

TTTAAGAGTG GATGCTCATA CAGCTTATTT TAATGGCAAT ATTTATCTGG GAAAATCCAC	60
GAATTTAAGA GTGAATGGCC ATAGCGCTCA TTTTAAAAAT ATTGATGCCA GTAAGAGCGA	120
TAACGGGGCTA AACACTACCA CTTTGGATTT CAGCGGCGTT ACAGATAAAG TCAATATCAA	180
CAAGCTCACT ACATCTGCCA CTAATGTGAA CATTAAAAAC TTTGACATTA AGGAATTAGT	240
GGTTACAACC CGAGTTCAGA GTTTTGGGCA ATACACTATT TTTGGCGAAA ATATAGGCGA	300
TAAGCTGCAC ATTGGTGTCTG TGAGTTTGCA AACGGGATAT AGCCCAGCCT ATTCTGGGGG	360
GCTTACTTTT AAAG	374

(2) INFORMATION FOR SEQ ID NO: 86:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 375 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 86:

TTTAAGAGTG GATGCTCATA CAGCTTATTT TAATGGCAAT ATTTATCTGG GAAAATCCAC	60
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GAATTTAAGA GTGAATGGCC ATAGCGCTCA TTTTAAAAAT ATTGATGCCA GTAAGAGCGA	120
TAACGGGCTA AACACTAGCT CTTTGGATTT CAGTGGCGTT ACAGACAAAG TCAATATCAA	180
CAAGCTCACT ACATCTGCCA CTAATGTGAA CGTTAAAAAC TTTGACATTA AGGAATTGGT	240
GGTTACAACC CGCGTTCAGA GTTTTGGGCA ATACACTATT TTTGGCGAAA ATATAGGCGA	300
TAAGTCTCGC ATTGGTGTCTG TTAGTTTGCA AACGGGATAT AGCCCGGCCT ATTCTGGGGG	360
CGTTACTTTT AAAAG	375

(2) INFORMATION FOR SEQ ID NO: 87:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 365 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 87:

GATGCTCATA CAGCTTATTT TAATGGCAAT ATTTATCTGG GAAAATCCAC GAATTTAAGA	60
GTGAATGGCC ATAGCGCTCA TTTTAAAAAT ATTGATGCCA GTAAGAGCGA TAACGGGCTA	120
AACACTAGCG CTTTGGATTT YAGCGGCGTT ACAGAYAAAG TCAATATCAA CAAGCTCACT	180
ACATCTGCCA CTAATGTGAA CGTTAAAAAC TTTGACATTA AGGAATTAGT GGTTACAACC	240
CGAGTTCAAA GTTTTGGGCA ATACACTATT TTTGGCGAAA ATATAGGCGA TAAGTCTCGC	300
ATTGGTGTCTG TTAGTTTGCA AACGGGATAT AGCCCGGCCT ATTCTGGGGG CGTTACTTTT	360
AAAAG	365

(2) INFORMATION FOR SEQ ID NO: 88:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 375 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 88:

TTTAAGAGGG GATGCTCATA CAGCTTATTT TAATGGCAAT ATTTATTTGG GAAAATCCAC	60
GAATTTAAGA GTGAATGGCC ATAGCGCTCA TTTTAAAAAT ATTGATGCCA GTAAGAGCGA	120
TAACGGGCTA AACACTAGCG YTTTGGATTT TAGCGGCGTT ACAGAYAAAG TCAATATCAA	180
CAAGCTCACT ACATCTGCCA CTAATGTGAA CRTTAAAAAC TTTGAYATTA AGGAATTGGT	240
GGTTACAACC CGAGTTCAAA GTTTTGGGCA ATACACTATT TTTGGCGAAA ATATAGGCGA	300
TMAGTCTCGC ATTGGTGTCTG TTAGTTTGCA AACGGGATAT AGCCCRGCCT ATTCTGGGGG	360
CGTTACTTTT AAAAG	375

(2) INFORMATION FOR SEQ ID NO: 89:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 375 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 89:

TTTAAGCGTG GATGCTCATA CAGCTTATTT TAATGGTAAT ATTTATCTGG GAAAATCCAC	60
GAATTTAAGA GTGAATGGCC ATAGCGCTCA TTTTAAAAAT ATTGATGCCA CAAAGAGCGA	120
TAACGGGCTA AACACTAGCG CTTTGGATTT CAGCGGCGTT ACAGATAAAG TCAATATCAA	180
CAAGCTCACT ACATCTGCCA CTAACGTGAA CATTAAAAAC TTTGACATTA AGGAATTGGT	240
GGTTACAACC CGAGTTCAAA GTTTTGGGCA ATACACTATT TTTGGCGAAA ATATAGGCGA	300
TAAGTCTCGC ATTGGTGTCTG TGAGTTTGCA AACGGGATAT AGCCCGGCCT ATTCTGGGGG	360
CGTTACTTTT AAAAG	375

(2) INFORMATION FOR SEQ ID NO: 90:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 375 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 90:

TTTAAGAGTG GATGCTCATA CAGCTTATTT TAATGGCAAT ATTTATCTGG GAAAATCCAC	60
GAATTTAAAA GTGAATGGCC ATAGCGCTCA TTTTAAAAAT ATTGATGCCA GTAAGAGCGA	120
TAATGGTCTA AACACTAGTG CTTTGGATTT GAGCGGCGTT ACAGACAAAG TCAATATCAA	180
CAAGCTCACT ACAGCTGCCA CTAATGTGAA CATTAAAAAC TTTGACATTA AGGAATTAGT	240
GGTTACGACC CGTGTTGAGA GTTTTGGGCA ATACACTATT TTTGGCGAAA ATATAGGAGA	300
TCAATCGCGC ATTGGTGTG TTAGTTTGCA AACTGGCTAT AGCCCGGCCT ATTCTGGGGG	360
CGTTACTTTT AAAAG	375

(2) INFORMATION FOR SEQ ID NO: 91:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 375 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 91:

CTTAAGAGTG GATGCTCATA CAGCTTATTT TAATGGCAAT ATTTATCTGG GAAAATCCAC	60
GAATTTAAGA GTGAATGGCC ATAGCGCTCA TTTTAAAAAT ATTGATGCTA GTAAGAGCGA	120
TAACGGGCTA AACACTAGCG CTTTGGATTT TAGCGGCGTT ACAGACAAAG TCAATATCAA	180
CAAGCTCACT ACATCTGCCA CTAATGTGAA CATTAAAAAC TTTGACATTA AGGAATTGGT	240

GGTTACAACC CGAGTTCAAA GTTTTGGGCA ATACACTATT TTTGGCGAAA ATATAGGCGA	300
TAAGTCTCGC ATTGGTGTCTG TGAGTTTGCA AACGGGATAT AGCCCGGCCT ATTCTGGGGG	360
CGTTACTTTT AAAAG	375

(2) INFORMATION FOR SEQ ID NO: 92:

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 449 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 92:

ATGACTAACG AAACCATTA CCAACAACCA CAAAGCGAAG CGGCTTTTAA CCCGCAGCAA	60
TTTATCAATA ATCTTCAAGT AGCTTTTCTT AAAGTTGATA ACGCTGTCGC TTCATACGAT	120
CCTGATCAAA AACCAATCGT TGATAAGAAC GATAGGGATA ATAGGCAAGC TTTTGATGGA	180
ATCTCGCAAT TAAGGGAAGA ATACTCCAAT AAAGCGATCA AAAATCCTAC CAAAAAGAAT	240
CAGTATTTTT CAGACTTTAT CAATAAGAGC AATGATTTAA TCAACAAAGA CGCTCTCATT	300
GATGTAGAAT CTTCCACAAA GAGCTTTCAG AAATTTGGGG ATCAGCGTTA CCGAATTTTC	360
ACAAGTTGGG TGTCCCATCA AAACGATCCG TCTAAAATCA ACACCCGATC GATCCGAAAT	420
TTTATGGAAA ATATCATACA ACCCCCTAT	449

(2) INFORMATION FOR SEQ ID NO: 93:

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 449 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 93:

ATGACTAACG AAACCATTAA CCAACAACCA CAAACCGAAG CGGCTTTTAA CCCGCAGCAA	60
TTTATCAATA ATCTTCAAGT AGCTTTTCTT AAAGTTGATA ACGCTGTCGC TTCATACGAT	120
CCTGATCAAA AACCAATCGT TGATAAGAAC GATAGGGATA ACAGGCAAGC TTTTGATGGA	180
ATCTCGCAAT TAAGGGAAGA ATACTCCAAT AAAGCGATCA AAAATCCTAC CAAAAAGAAT	240
CAGTATTTTT CAGACTTTAT CAATAAGAGC AATGATTTAA TCAACAAAGA CGCTCTCATT	300
GATGTAGAAT CTTCCACAAA GAGCTTTCAG AAATTTGGGG ATCAGCGTTA CCGAATTTTC	360
ACAAGTTGGG TGTCCCATCA AAACGATCCG TCTAAAATCA ACACCCGATC GATCCGAAAT	420
TTTATGGAAA ATATCATACA ACCCCCTAT	449

(2) INFORMATION FOR SEQ ID NO: 94:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 449 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 94:

ATGGCTAACG AAACATTAA CCAACAACCA CAAACCGAAG CGGCTTTTAA CCCGCAGCAA	60
TTTATCAATA ATCTTCAAGT AGCTTTTCTT AAAGTTGATA ACGCTGTCGC TTCATACGAT	120
CCTGATCAAA AACCAATCGT TGATAAGAAC GATAGGGATA ACAGGCAAGC TTTTGATGGA	180
ATCTCGCAAT TAAGGGAAGA ATACTCCAAT AAAGCGATCA AAAATCCTAC CAAAAAGAAT	240
CAGTATTTTT CAGACTTTAT CAATAAGAGC AATGATTTAA TCAACAAAGA CGCTCTCATT	300
GATGTAGAAT CTTCCACAAA GAGCTTTCAG AAATTTGGGG ATCAGCGTTA CCGAATTTTC	360
ACAAGTTGGG TGTCCCATCA AAACGATCCG TCTAAAATCA ACACCCGATC GATCCGAAAT	420
TTTATGGAAA ATATCATACA ACCCCCTAT	449

(2) INFORMATION FOR SEQ ID NO: 95:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 449 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 95:

ATGACTAACG AAACCATTAA CCAACAACCA CAAACCGAAG CGGCTTTTAA CCCGCAGCAA	60
TTTATCAATA ATCTTCAAGT GGCTTTTCTT AAAGTTGATA ACGCTGTCGC TTCATACGAT	120
CCTGATCAAA AACCAATCGT TGATAAGAAC GATAGGGATA ATAGGCAAGC TTTTGATGGA	180
ATCTCGCAAT TAAGGGAAGA ATACTCCAAT AAAGCGATCA AAAATCCTAC CAAAAAGAAT	240
CAGTATTTTT CAGACTTTAT CAATAAGAGC AATGATTAA TCAACAAAGA CGCTCTCATT	300
GATGTAGAAT CTTCACAAA GAGCTTTCAG AAATTTTGGG ATCAGCGTTA CCGAATTTTC	360
ACAAGTTGGG TGTCCCATCA AAACGATCCG TCTAAAATCA ACACCCGATC GATCCGAAAT	420
TTCATGGAAA ATATCATACA ACCCCCTAT	449

(2) INFORMATION FOR SEQ ID NO: 96:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 449 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 96:

ATGACTAACG AAACCATTAA CCAACAACCA CAAACCGAAG CGGCTTTTAA CCCGCAGCAA	60
TTTATCAATA ATCTTCAAGT GGCTTTTCTT AAAGTTGATA ACGCTGTCGC TTCATACGAT	120
CCTGATCAAA AACCAATCGT TGATAAGAAC GATAGGGATA ACAGGCAAGC TTTTGATGGA	180

ATCTCGCAAT TAAGGGAAGA ATACTCCAAT AAAGCGATCA AAAATCCTAC CAAAAGAAT	240
CAGTATTTTT CAAACTTTAT CAATAAGAGC AATGATCTAA TCAACAAAGA CAATCTCATT	300
GATGTAGAAT CTTCCAAAAA GAGCTTTCAG AAATTTGGGG ATCAGCGTTA CCGAATTTTC	360
ACAAGTTGGG TGTCCCATCA AAACGATCCG TCTAAATCA ACACCCGATC GATCCGAAAT	420
TTTATGGAAA ATATCATACA ACCCCCTAT	449

(2) INFORMATION FOR SEQ ID NO: 97:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 449 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 97:

ATGACTAACG AAACCATTAA CCAACAACCA CAAACCGAAG CGGCTTTTAA CCCGCAGCAA	60
TTTATCAATA ATCTTCAAGT AGCTTTTCTT AAAGTTGATA ACGCTGTCGC TTCATACGAT	120
CCTGACCAAA AACCAATCGT TGATAAGAAC GATAGGGATA ACAGGCAAGC TTTTGATAGA	180
ATCTCACAAT TAAGGGAGGA ATACTCCAAT AAAGCGATCA AAAATCCTAC CAAAAGAAT	240
CAGTATTTTT CAGACTTTAT CGATAAGAGC AACGATTTAA TCAACAAAGA CGCTCTCATT	300
GATGTAGAAT CTTCCACAAA GAGCTTTCAG AAATTTGGGG ATCAGCGTTA CCGAATTTTC	360
ACAAGTTGGG TGTCCCATCA AAACGATCCG TCTAAATCA ACACCCGATC GATCCGAAAT	420
TTTATGGAAA ATATCATACA ACCCCCTAT	449

(2) INFORMATION FOR SEQ ID NO: 98:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 449 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 98:

ATGACTAACG AAACCATTAA CCAACAACCA CAAACCGAAG CGGCTTTTAA CCCGCAGCAA	60
TTTATCAATA ATCTTCAAGT GGCTTTTCTT AAAGTTGATA ACGCTGTCGC TTCATACGAT	120
CCTGATCAAA AACCAATTAT TGATAAGAAC GATAGGGATA ACAGGCAAGC TTTTGATGGA	180
ATCTCGCAAT TAAGGGAAGA ATATTCCAAT AAAGCGATCA AAAATCCTAC CAAAAAGAAT	240
CAGTATTTTT CAGACTTTAT CGATAAGAGC AATGATTTAA TCAACAAAGA CAATCTCATT	300
GATGTAGAAT CTTCCACAAA GAGCTTTCAG AAATTTGGGG ATCAGCGTTA CCGAATTTTC	360
ACAAGTTGGG TGTCCCATCA AAACGATCCG TCTAAAATCA ACACCCGATC GATCCGAAAT	420
TTTATGGAAA ATATCATACA ACCCCCTAT	449

(2) INFORMATION FOR SEQ ID NO: 99:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 449 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 99:

ATGACTAACG AAACCATTAA CCAACAACCA CAAACCGAAG CGGCTTTTAA CCCGCAGCAA	60
TTTATCAATA ATCTTCAAGT AGCTTTTCTT AAAGTTGATA ATGCTGTCGC TTCATACGAT	120
TCTGATCAAA AACCAATCAT TGATAAGAAC GATAGGGATA ACAGGCAAGC TTTTGATAGA	180
ATCTCGCAAT TAAGGGAAGA ATACTCCAAT AAAGCGATCA AAAATCCTAC CAAAAAGAAT	240
CAGTATTTTT CAGACTTTAT CGATAAGAGC AACGATTTAA TCAACAAAGA CAATCTCATT	300
GATGTAGAAT CTTCCACAAA GAGCTTTCAG AAATTTGGGG ATCAGCGTTA CCGAATTTTC	360
ACAAGTTGGG TGTCCCATCA AAATGATCCG TCTAAAATCA ACACCCGATC GATCCGAAAT	420
TTTATGGAAA ATATCATACA ACCCCCTAT	449

(2) INFORMATION FOR SEQ ID NO: 100:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 449 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 100:

ATGACTAACG AACTATTGA CCAACAACCA CAAACCGAAG CGGCTTTTAA CCCGCAGCAA	60
TTTATTAATA ATCTTCAGGT AGCTTTTCTT AAGCTTGATA ACGCTGTCGC TTCATTGAT	120
CCTGATCAAA AACCAATCGT TGATAAGAAT GATAGGGATA ACAGGCAAGC TTTTGATGGA	180
ATCTCGCAAT TAAGGGAAGA ATACTCCAAT AAAGCGATCA AAAATCCTAC CAAAAAGAAT	240
CAGTATTTTT CAGACTTTAT CAATAAGAGC AATGATTTAA TCAACAAAGA CGCTCTCATT	300
GATGTAGAAT CTTCCACAAA GAGCTTTCAG AAATTTGGGG ATCAGCGTTA CCGAATTTTC	360
ACAAGTTGGG TGTCCCATCA AAACGATCCG TCTAAAATCA ACACCCGATC GATCCAAAAT	420
TTTATGGAAA ATATCATACA ACCCCCTAT	449

(2) INFORMATION FOR SEQ ID NO: 101:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 449 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 101:

ATGACTAACG AACTATTGA CCAACAACCA CAAACTGAAG CGGCTTTTAA CCCGCAGCAA	60
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TTTATCAATA ATCTTCAAGT GGCTTTTCTT AAGCTTGATA ACGCTGTCGC TTCATTTGAT	120
CCTGATCAAA AACCAATCGT TGATAAGAAC GATAGGGATA ACAGGCAAGC TTTTGATGGA	180
ATCTCGCAAT TAAGGGAAGA ATACTCCAAT AAAGCGATCA AAAATCCTAC CAAAAAGAAT	240
CAGTATTTTT CAGACTTTAT CAATAAGAGC AATGATTTAA TCAACAAAGA CGCTCTCATT	300
GATGTAGAAT CTTCCACAAA GAGCTTTCAG AAATTTGGGG ATCAGCGTTA CCGAATTTTC	360
ACAAGTTGGG TGTCCCATCA AAACGATCCG TCTAAAATCA ACACCCGATC GATCCGAAAT	420
TTTATGGAAA ATATCATACA ACCCCCTAT	449

(2) INFORMATION FOR SEQ ID NO: 102:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 449 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 102:

ATGACTAACG AAACATTTAA CCAACAGCCA CAAACCGAAG CGGCTTTTAA CCCGCAGCAA	60
TTTATCAATA ATCTTCAAGT AGCTTTTCTT AAGCTTGATA ACGCTGTCGC TTCATTTGAT	120
CCTGATCAAA AACCAATCGT TGATAAGAAC GATAGGGATA ATAGGCAGGC TTTTGATGGA	180
ATCTCGCAAT TAAGGGAAGA ATACTCCAAT AAAGCGATCA AAAATCCTAC CAAAAAGAAT	240
CAGTATTTTT CAGACTTTAT CAATAAGAGC AATGATTTAA TCAACAAAGA CAATCTCATT	300
GATGTAGAAT CTTCCACAAA GAGCTTTCAG AAATTTGGGG ATCAGCGTTA CCGAATTTTC	360
ACAAGTTGGG TGTCCCATCA AAACGATCCG TCTAAAATCA ACACCCGATC GATCCGAAAT	420
TTTATGGAAA ATATCATACA ACCCCCTAT	449

(2) INFORMATION FOR SEQ ID NO: 103:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 449 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 103:

ATGACTAACG AAACCATTAA CCAACAACCA CAAACCGAAG CGGCTTTTAA CCCGCAGCAA	60
TTTATCAATA ATCTTCAAGT GGCTTTTCTT AAGCTTGATA ATGCTGTTGC TTCATTTGAT	120
CCTGATCAAA AACCAATCGT TGATAAGAAC GATAGGGATA ACAGGCAAGC TTTTGATGGA	180
ATCTCGCAAT TAAGGGAAGA ATACTCCAAT AAAGCGATCA AAAATCCTAC CAAAAAGAAT	240
CAGTATTTTT CAGACTTTAT CGATAAGAGC AACGATTTAA TCAACAAAGA CAATCTCATT	300
GATGTAGAAT CTTCCACAAA GAGCTTTCAG AAATTTGGGG ATCAGCGTTA CCGAATTTTC	360
ACAAGTTGGG TGTCCCATCA AAACGATCCG TCTAAAATCA ACACCCGATC GATCCGAAAT	420
TTTATGGAAA ATATCATACA ACCCCCTAT	449

(2) INFORMATION FOR SEQ ID NO: 104:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 449 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 104:

ATGACTAACG AAACCATTAA CCAACAACCA CAAACCGAAG CGGCTTTTAA CCCGCAGCAA	60
TTTATCAATA ATCTTCAAGT GGCTTTTCTT AAAGTTGATA ACGCTGTTCG TTCATACGAT	120
CCTGATCAAA AACCAATCGT TGATAAGAAC GATAGGGATA ATAGGCAAGC TTTTGATGGA	180
ATCTCGCAAT TAAGGGAAGA ATACTCCAAT AAAGCGATCA AAAATCCTAC CAAAAAGAAT	240
CAGTATTTTT CAGACTTTAT CAATAAGAGC AATGATTTAA TCAACAAAGA CGCTCTCATT	300
GATGTAGAAT CTTCCACAAA GAGCTTTCAG AAATTTTGGG ATCAGCGTTA CCGAATTTTC	360

ACAAGTTGGG TGTCCCATCA AAACGATCCG TCTAAATCA ACACCCGATC GATCCGAAAT	420
TTCATGGAAA ATATCATACA ACCCCCTAT	449

(2) INFORMATION FOR SEQ ID NO: 105:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 449 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 105:

ATGACTAACG AAACATTTGA TCAACAACCA CGAACCGAAG CGGCTTTTAA CCCGCAGCAA	60
TTTATCAATA ATCTTCAAGT AGCTTTTCTT AAAGTTGATA ACGTTGTCGC TTCATTTGAT	120
CCTAATCAAA AACCAATCGT TGATAAGAAC GATAGGGATA ACAGGCAAGC TTTTGATGGA	180
ATCTCGCAAT TAAGGGAAGA ATACTCCAAT AAAGCGATCA AAAATCCTGC CAAAAAGAAT	240
CAGTATTTTT CAGACTTTAT CAATAAGAGC AATGATCTAA TCAACAAAGA CAATCTCATT	300
GATGTAGAAT CTTCCACAAA GAGCTTTCAG AAATTTGGGG ATCAGCGTTA CCAAATTTTC	360
ACAAGTTGGG TGTCCCATCA AAACGATCCG TCTAAATCA ACACCCGATC GATCCGAAAT	420
TTTATGGAAA ATATCATACA ACCCCCTAT	449

(2) INFORMATION FOR SEQ ID NO: 106:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 449 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 106:

ATGACTAACG AAACCATTAA CCAACAACCA CAAACCGAAG CGGCTTTTAA CCCGCAGCAA	60
TTTATCAATA ATCTTCAAGT GGCTTTTATT AAAGTTGATA ATGTTGTCGC TTCATTTGAT	120
CCTGATCAAA AACCAATCGT TGATAAGAAT GATAGGGATA ATAGGCAAGC TTTTGAGAAA	180
ATCTCGCAGC TAAGGGAGGA ATTCGCTAAT AAAGCGATCA AAAATCCTGC CAAAAAGAAT	240
CAGTATTTTT CAAGCTTTAT CAGTAAGAGC AGTGATTTAA TCAACAAAGA CAGTCTCATT	300
GATACAGGTT CTTCCATAAA GAGCTTTCAG AAATTGGGA CTCAGCGTTA CCAAATTTTT	360
ATGAATTGGG TGTCCCATCA AAAAGATCCA TCTAAATCA ACACCCAAAA AATCCGAGGT	420
TTTATGGAAA ATATCATACA ACCCCCTAT	449

(2) INFORMATION FOR SEQ ID NO: 107:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 464 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 107:

ATGACTAACG AAACATTGA TCAACAAGA ACACCAGACC AAACACAAAG CCAAACAGCT	60
TTTGATCCGC AACAATTTAT CAATAATATT CAAGTGGCTT TTCTTAAAGT TGATAACGCT	120
GTCGCTTCAT TTGATCCTGA TCAAAAACCA ATCGTTGATA AGAATGATAG GGATAACAGG	180
CAAGCTTTTG AGAAAATCTC GCAGCTAAGG GAGGAATTCG CTAATAAAGC GATCAAAAAT	240
CCTGCCAAAA AGAATCAGTA TTTTCAAGC TTTATCAGTA AGAGCAGTGA TTTAGTCAAC	300
AAAGACAGTC TCATTGATAC AGGTTCTTCC ATAAAGAGCT TTCAGAAATT TGGGACTCAG	360
CGTTACCAAA TTTTATGAA TTGGGTGTCC CATCAAAAAG ATCCATCTAA AATCAACACC	420
CAAAAATCC AAGATTTTAT GGAAAATATC ATACAACCCC CTAT	464

(2) INFORMATION FOR SEQ ID NO: 108:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 464 base pairs

(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 108:

ATGACTAATG AAACCATTGA TCAAACAACA ACACCAGATC AAACACCAA TCAAACAGAT	60
TTTGTTCGCG AACGATTTAT CAATAATCTT CAAGTAGCTT TTATTAAAGT TGATAACGCT	120
GTCTCTTCAT TTGATCCTGA TCAAAAACCA ATCGTTGATA AGAATGATAG GGATAACAGG	180
CAAGCTTTTG AGAAAATCTC GCAACTAAGG GAAGAATACG CCAATAAAGC GATCAAAAAT	240
CCTGCCAAAA AGAATCAGTA TTTTTCAGAC TTTATCAATA AGAGCAATGA TTTGATCAAC	300
AAAGACAATC TCATTGCTGT AGATTCTTCC GTAGAGACCT TTAAGAAATT TGGGGATCAG	360
CGTTACCAA TTTTACGAA TTGGGTGTCC CTTCAAAAAG ATCCGTCTAA AATCAACACC	420
CGACAAATCC GAAATTTTAT GGAAAATATC ATACAACCCC CTAT	464

(2) INFORMATION FOR SEQ ID NO: 109:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 464 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 109:

ATGACTAACG AAACCATTGA TCAAACAACA ACACCAGATC AAACACCAA CCAAACGGAT	60
TTTGTTCGCG AACGATTTAT CAATAATCTT CAAGTAGCTT TCCTTAAAGT TGATAGCGCT	120
GTGCTTCAT TTGATCCTGA TCAAAAACCA ATCGTTGATA AGAATGATAG GGATAACAGG	180
CAAGCTTTTG AGAAAATCTC GCAACTAAGG GAAGAATACG CCAATAAAGC GATCAAAAAT	240

CCTGCCAAAA AGAATCAGTA TTTTTCAGAC TTTATCAATA AGAGCAATGA TTTGATCAAC	300
AAAGACAATC TCATTGCTGT AGATTCTTCT GTAGAGAGCT TTAAGAAATT TGGGGATCAG	360
CGTTACCAAA TTTTACGAG TTGGGTGTCC CTTCAAAAAG ATCCGTCTCA AATCAACACC	420
CGACAAATCC GAAATTTTAT GGAAAATATC ATACAACCCC CTAT	464

(2) INFORMATION FOR SEQ ID NO: 110:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 464 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 110:

ATGACTAACG AAACCATTGA TCAAACAACA ACACCAGATC AAACACCAAG CCAAACAGAT	60
TTTGTTCGCG AACGATTTAT CAATAATCTT CAAGTAGCTT TTCTTAAAGT TGATAACGCT	120
GTCGCTTCAT TTGATCCTGA TCAAAAACCA ATCGTTGATA AGAATGATAG GGATAACAGG	180
CAAGCTTTTG AGAAAATCTC GCAACTAAGG GAAGAATACG CCAATAAAGC GATCAAAAAT	240
CCTGCCAAAA AGAATCAGTA TTTTTCAGAC TTTATCAATA AGACCAATGA TTTGATCAAC	300
AAAGACAATC TCATTGCTGT AGATTCTTCC GTAGATAGCT TTAAGAAATT TGGGGATCAG	360
CGTTACCAAA TTTTACGAG TTGGGTGTCC CTTCAAAAAG ATCCGTCTAA AATCAACACC	420
CAACAAATCC GAAATTTTAT GGAAAATATC ATACAACCCC CTAT	464

(2) INFORMATION FOR SEQ ID NO: 111:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 464 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 111:

ATGACTAACG AAACCATTGA TCAAACAACA ACACCAGATC AAACACCAAA TCAAACAGAT	60
TTTGTTCGCG AACGATTTAT CAATAATCTT CAAGTAGCTT TTATTAAAGT TGATGACGCT	120
GTCGCTTCAT TTGATCCCGA TCAAAAACCA ATCGTTGATA AGAATGATAG GGATAACAGG	180
CAAGCTTTTG AGAAAATCTC GCAACTAAGG GAAGAATACG CCAATAAAGC GATCAAAAAT	240
CCCACCAAAA AGAATCAGTA TTTTTCAGAC TTTATCAATA AGACCAATGA TTTGATCAAC	300
AAAGACAATC TCATTGCTGT AGATTCTTCC GTAGAGAGCT TTAAGAAATT TGGGGATCAG	360
CGTTACCAAA TTTTACGAG TTGGGTGTCC CTTCAAAAAG ATCCGTCTAA AATCAACACC	420
CAACAAATCC GAAATTTTAT GGAAAATATC ATACAACCCC CTAT	464

(2) INFORMATION FOR SEQ ID NO: 112:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 464 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 112:

ATGACTAACG AAACCATTGA TCAAACAACA ACACCAGATC AAACACCAAA TCAAACAGAT	60
TTTGTTCGCG AACGATTTAT CAATAATCTT CAAGTAGCTT TTATTAAAGT TGATAACGCT	120
GTTGCTTCAT TTGATCCCGA TCAAAAACCA ATCGTTGATA AGAATGATAG GGATAACAGG	180
CAAGCTTTTG AGAAAATCTC GCAACTAAGG GAAGAATACG CCAATAAAGC GATCAAAAAT	240
CCTGCCAAAA AGAATCAGTA TTTTTCAGAC TTTATCAATA AGAGCAATGA TTTGATCAAC	300
AAAGACAATC TCATTGCTGT AGATTCTTCC GTAGATAGCT TTAAGAAATT TGGGGATCAG	360
CGTTACCAAA TTTTACGAG TTGGGTGTCC CTTCAAAAAG ATCCGTCTCA AATCAACACC	420
CAACAAATCC AAAATTTTAT GGAAAATATC ATACAACCCC CTAT	464

(2) INFORMATION FOR SEQ ID NO: 113:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 464 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 113:

ATGACTAACG AAACCATTGA TCAAACAACA ACACCAGATC AAACACTAAA CCAAACGGAT	60
TTTGTTC CGC AACGATTTAT CAATAATCTT CAAGTAGCTT TTATTAAAGT TGATAACGCT	120
GTCGCTTTAT TTGATCCCGA TCAAAAACCA ATCGTTGATA AGAATGATAG GGATAACAGG	180
CAAGCTTTTG AGAAAATCTC GCAACTAAGG GAAGAATACG CCAATAAAGC GATCAAAAAT	240
CCCACCAAAA AGAATCAGTA TTTTTCAGAC TTTATCAATA AGAGCAATGA TTTGATCAAC	300
AAAGACAATC TCATTGCTGT AGATTCTTCC GTAGATAGCT TTAAGAAATT TGGGGATCAG	360
CGTTACCAAA TTTTACGAG TTGGGTGTCC CTTCAAAAAG ATCCGTCTCA AATCAACACC	420
CGACAAATCC GAAATTTTAT GGAAAATATC ATACAACCCC CTAT	464

(2) INFORMATION FOR SEQ ID NO: 114:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 464 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 114:

ATGACTAACG AAACCATTGA TCAAACAATA ACACCAGATC AAACACCAAA CCAAACGGAT	60
TTTGTTC CGC AACGATTTAT CAATAATCTT CAAGTAGCTT TTATCAAAGT TGATAACGCT	120

GTCGCTTCAT TTGATCCTGA TCAAAAACCA ATCGTTGATA AGAATGATAG GGATAACAGG	180
CAAGCTTTTG AGAAAATCTC GCAATTAAGG GAAGAATAcG CCAATAAAGC GATCAAAAAT	240
CCTGCCAAAA AGAATCAGTA TTTTTTAGAC TTTATCAATA AGAGCAATGA TTTGATCAAC	300
AAAGACAATC TCATTGCTGT AGATTCTTCC GTAGATAGCT TTAAGAAATT TGGGGATCAG	360
CGTTACCAAA TTTTACGAG TTGGGTGTCC CTTCAAAAAG ATCCGTCTAA AATCAACACC	420
CAACAAATCC GAAATTTTAT GGAAAATATC ATACAACCCC CTAT	464

(2) INFORMATION FOR SEQ ID NO: 115:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 132 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 115:

CCCTATTATC TCTCTCGCTC TAGTGGGGGT GTTAATGGGT ACCGAACTAG GGGCTAACAC	60
GCCAAACGAT CCCATACACA GCGAGAGTCG CGCCTTTTTT ACAACCGTGA TCATTCCAGC	120
CATTGTTGGG GG	132

(2) INFORMATION FOR SEQ ID NO: 116:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 132 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 116:

CCCTATTATC TCTCTCGCTC TAGTGGGGGT GTTAATGGGT ACCGAACTAG GGGCTAACAC	60
GCCAAACGAT CCCATACACA GCGAGAGTCG CGCCTTTTTC ACAACCGTGA TCATTCCAGC	120
CATTGTTGGA GG	132

(2) INFORMATION FOR SEQ ID NO: 117:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 132 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 117:

CCCTATTATC TCTCTCGCTC TAGTGGGGGT GTTAATGGGC ACCGAACTAG GGGCTAATAC	60
GCCAAACGAT CCCATACACA GCGAGAGTCG CGCCTTTTTC ACAACCGTGA TCATTCCAGC	120
CATTGTTGGG GG	132

(2) INFORMATION FOR SEQ ID NO: 118:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 132 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 118:

CCCTATTATC TCTCTCGCTC TAGTGGGGGT GTTAATGGGC ACCGAACTAG GGGCTAATAC	60
GCCAAACGAT CCCATACACA GCGAGAGTCG CGCCTTTTTC ACAACCGTGA TCATTCCAGC	120
CATTGTTGGG GG	132

(2) INFORMATION FOR SEQ ID NO: 119:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 132 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 119:

CCCTATTATC TCTCTCGCTC TAGTGGGGGT GTTAATGGGC ACTGAACTAG GGGCTAACAC	60
GCCAAACGAT CCCATACACA GCGAGAGTCG CGCCTTTTTT ACAACCGTGA TCATTCCAGC	120
CATTGTTGGG GG	132

(2) INFORMATION FOR SEQ ID NO: 120:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 132 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 120:

CCCTATTATC TCTCTCGCTC TAGTGGGGGT GTTAATGGGC ACCGAACTAG GGGCTAACAC	60
GCCAAATGAT CCCATACACA GCGAGAGTCG CGCCTTTTTT ACAACYGTGA TCATTCCAGC	120
CATTGTTGGG GG	132

(2) INFORMATION FOR SEQ ID NO: 121:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 132 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 121:

CCCTATTATC TCTCTCGCTT TAGTGGGGGT RTAATGGGC ACCGAAGTAG GGGCTAACAC	60
GCCAAACGAT CCCATACACA GCGAGAGTCG CGCCTTTTTC ACAACCGTGA TCATTCCAGC	120
CATTGTTGGG GG	132

(2) INFORMATION FOR SEQ ID NO: 122:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 132 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 122:

CCCTATTATT TCTCTCGCTT TAGTGGGGGT GTTAATGGGC ACCGAAGTAG GGGCTAACAC	60
GCCAAACGAT CCCATACACA GCGAGAGTCG CGCCTTTTTT ACAACCGTGA TCATTCCAGC	120
CATTGTTGGG GG	132

(2) INFORMATION FOR SEQ ID NO: 123:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 132 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 123:

```
CCCTATTATC TCTCTCGCTC TAGTGGGGGC ATTAATGAGT ACCGAACTAG GGGCTAACAC      60
GCCAAATGAT CCCATACACA GCGAGAGTCG CGCCTTTTTT ACAACCGTGA TCATTCCAGC      120
CATTGTTGGG GG                                                              132
```

(2) INFORMATION FOR SEQ ID NO: 124:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 132 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 124:

```
CCCTATTATC TCTCTCGCTT TAGTGGGGGT ATTAATAGGC ACCGAACTAG GGGCTAATAC      60
GCCAAATGAT CCCATACACA GCGAGAGTCG TGCTTTTTTC ACAACCGTGA TCATTCCAGC      120
CATTGTTGGG GG                                                              132
```

(2) INFORMATION FOR SEQ ID NO: 125:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 132 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 125:

```
CCCTATTATC TCTCTCGCTC TAGTGGGGGT GTTAATAAGC ACCGAACTAG GGGCTAACAC      60
GCCAAATGAT CCCATACACA GCGAGAGTCG CGCCTTTTTT ACAACGGGGA TCATTCCAGC      120
```


CATTGTTGGG GG

132

(2) INFORMATION FOR SEQ ID NO: 126:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 126:

CCCTTTAGTT TCTCTCGCTT TAGTGGGGTT ATTGGTCAGC ATCACACCAC AAAAAAGTCA 60

TGCTGCCTTC TTTACAACCG TGATCATTC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 127:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 127:

CCCTTTAGTT TCTCTCGCTT TAGTGGGGTT ATTGGTCAGC ATCACACCAC AAAAAAGTCA 60

TGCTGCCTTC TTTACAACCG TGATCATTC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 128:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 128:

CCCTTTAGTT TCTCTCGCTT TAGTGGGGTT ATTGGTCAGC ATCACACCGC AAAAAAGTCA 60

TGCTGCCTTC TTTACAACCG TGATCATTC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 129:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 105 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 129:

CCCTCTAGTT TCTCTCGCTT TAGTGGGGTT ATTGGTCAGC ATCACACCGC AAAAAAGTCA 60

TGCCGCCTTC TTTACAACCG TGATCATTC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 130:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 105 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 130:

CCCTCTAGTT TCTCTCGCTT TAGTGGGGTT ATTGGTCAGC ATCACACCGC AAAAAAGTCA 60

TGCCGCCTTC TTTACAACCG TGATCATTC ADCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 131:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 131:

CCCTTTAGTT TCTCTCGCTT TAGTGGGGTT ATTGGTCAGC ATCACACCGC AAAAAAGTCA 60

TGCCGCCTTC TTTACAACCG TGATCATTC GGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 132:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 132:

CCCTTTAGTT TCTCTCGCTT TAGTGGGGTT ATTGGTCAGC ATCACACCGC AAAAAAGTCA 60

TGCCGCCTTC TTTACAACCG TGATCATTC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 133:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 133:

CCCTTTAGTT TCTCTCGCTT TAGTGGGGCT ATTGGTCAGC ATCACACCGC AAAAAAGTCA 60

TGCCGCCTTC TTTACAACCG TGATCATTC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 134:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 105 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 134:

CCCTTTAGTT TCTCTCGCTT TAGTGGGGTT ATTGGTCAGC ATCACACCGC AAAAAAGTCA 60

TGCCGCCTTT TTTACAACCG TGATCATTC GGCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 135:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 105 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 135:

CCCTTTAGTT TCTCTCGCTT TAGTGGGGCT ATTGGTCAGC ATCACACCGC AAAAAAGTCA 60
 TGCCGCCTTC TTTACAACCG TGATTATCCC GGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 136:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 136:

CCCTTTAGTT TCTCTCGCTT TAGTGGGGCT ATTGGTCAGC ATCACACCAC AAAAAAGTCA 60
 TGCCGCCTTT TTTACAACCG TGATCATTCC AGCCATTGTT GGGAG 105

(2) INFORMATION FOR SEQ ID NO: 137:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 137:

CCCTTTAGTT TCTCTCGCTT TAGTGGGGCT ATTGGTCAGC ATCACACCAC AAAAAAGTCA 60
 TGCCGCCTTT TTTACAACCG TGATCATTCC AGCCATTGTT GGAGG 105

(2) INFORMATION FOR SEQ ID NO: 138:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 138:

CCCTCTGGTA TCTCTCGCTT TAGTGGGGCT ATTGGTCAGC ATCACACCGC AAAAAAGTCA	60
TGCCGCCTTC TTTACAACCG TGAATATTCC AGCMATTGTT GGGGG	105

(2) INFORMATION FOR SEQ ID NO: 139:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 139:

CCCTCTGGTT TCTCTCGCTT TAGTGGGGCT ATTGGTCAGC ATCACACCGC AAAAAAGTCA	60
TGCCGCCTTT TTCACAACCG TGATCATTCC GGCCATTGTT GGGGG	105

(2) INFORMATION FOR SEQ ID NO: 140:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 140:

CCCTCTGGTT TCTCTCGCTT TAGTGGGGCT ATTGGTCAGC ATCACACCGC AAAAAAGTCA 60
TGCCGCCTTC TTTACAACCG TGATCATTC AGCMATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 141:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 141:

CCCTCTGGTT TCTCTCGCTT TAGTGGGGCT ATTGGTCAGC ATCACACCGC AAAAAAGTCA 60
TGCCGCCTTC TTTACAACCG TGATCATTC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 142:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 142:

CCCTCTGGTT TCTCTCGCTT TAGTGGGGCT ATTGGTCAGC ATCACACCAC AAAAAAGTCA 60
TGCCGCCTTC TTTACAACCG TGATCATTC AGCHATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 143:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid

(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)
(iii) HYPOTHETICAL: NO
(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 143:

CCCTCTGGTT TCTCTCGCTT TAGTGGGGCT ATTGGTCAGC ATCACACCAC AAAAAAGTCA 60

TGCCGCCTTC TTTACAACCG TGATCATTC AGCYATTGTT GGAGG 105

(2) INFORMATION FOR SEQ ID NO: 144:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 105 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)
(iii) HYPOTHETICAL: NO
(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 144:

CCCTCTGGTT TCTCTCGCTT TAGTGGGGTT ATTGGTCAGC ATCACACCGC AAAAAAGTCA 60

TGCCGCCTTC TTTACAACCG TGATCATTC AGCCATTGTT GGAGG 105

(2) INFORMATION FOR SEQ ID NO: 145:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 105 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)
(iii) HYPOTHETICAL: NO
(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 145:

CCCTTTAGTT TCTCTCGCTT TAGTGGGGTT ATTGGTTAGC ATCACACCGC AAAAAAGTCA 60
TGCCGCCTTT TTCACAACCG TGATTATTCC AGCCATTGGT TGGGG 105

(2) INFORMATION FOR SEQ ID NO: 146:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 146:

CCCTCTGGTT TCTCTCGCTT TAGTGGGGCT ATTGGTCAGC ATCACACCAC AAAAAAGTCA 60
TGCCGCCTTC TTTACAACCG TGATCATTC AGCCATCGTT CCCCC 105

(2) INFORMATION FOR SEQ ID NO: 147:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 147:

CCCTTTAGTT TCTCCTGTTT TAGCAGGAGC GTTGATTAGC TCCATACCGC AACAAAGTCA 60
TGCCGCCTTT TTCACAACCG TGATCATTC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 148:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs

(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 148:

CCCTTTAGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGT GCCATACCGC AAGAAAGTCA 60

TGCCGCCTTT TTTACAACCG TAATCATTC CAGCTATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 149:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 105 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 149:

CCCTTTAGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC GCCATACCGC AAGAGAGTCA 60

TGCCGCCTTT TTCACGACCG TGATCATTC CAGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 150:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 105 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 150:

```
CCCTTTAGTT TCTCTCGTTT TAGCAGGAGC GTTATTAGC GCCATACCGC AAGAGAGTCA      60
TGCCGCCTTT TTCACGACCG TGATCATTCC AGCCATTGTT GGGGG                      105
```

(2) INFORMATION FOR SEQ ID NO: 151:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 151:

```
CCCTTTAGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC GCCATACCGC AAGAAAGTTA      60
TGCCGCCTTT TTCACGACCG TGATCATTCC AGCCATTGTT GGGGG                      105
```

(2) INFORMATION FOR SEQ ID NO: 152:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 152:

```
CCCTTTAGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC GCCATACCGC AACAAAGTCA      60
TGCCGCCTTT TTCACAACCG TGATCATTCC AGCCATTGTT GGGGG                      105
```

(2) INFORMATION FOR SEQ ID NO: 153:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 153:

```
CCCTATTATT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC TCCATACCGC AAGAAAGTCA      60
TGCCGCCTTT TTCACGACCG TGATCATTCC AGCCATTGTT GGGGG                      105
```

(2) INFORMATION FOR SEQ ID NO: 154:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 154:

```
CCCTTTAGTT TCTCTCGTTT TAGCAGNAGC GTTNATTAGT GCCATACCGC AANANAGTCA      60
TGCCGCCTTT TTCACNACCG TNATCATTCC AGCCATTGTT GGGGG                      105
```

(2) INFORMATION FOR SEQ ID NO: 155:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 155:

CCCTTTGGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC GCCATACCGC AACAAAGTCA 60
TGCCGCCTTT TTCACAACCG TGATCATTC AGCCATTGTG GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 156:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 156:

CCCTTTAGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC GCCATACCGC AACAAAGTCA 60
TGCCGCCTTT TTCACAACCG TGATCATTC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 157:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 157:

CCCTTTAGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC GCCATACCGC AACAAAGTCA 60
TGCCGCCTTT TTCACAACCG TGATCATTC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 158:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 158:

CCCTTTAGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC TCCATACCGC AACAAAGTCA 60

TGCCGCCTTT TTCACAACCG TGATCATTC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 159:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 159:

CCCTTTAGTT TCTCTTGT TT TAGCAGGAGC GTTGATTAGC GCCATACCGC AACAAAGTCA 60

TGCCGCCTTT TTCACGACCG TGATCATTC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 160:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 160:

CCCTTTAGTT TCTCTTGTTT TAGCAGGAGC GTTGATTAGC GCCATACCGC AACAAAGTCA 60
TGCCGCCTTT TTCACGACCG TGATCATTC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 161:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 105 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 161:

CCCTTTGGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC GCCATACCGC AACAAAGTCA 60
TGCCGCCTTT TTCACAACCG TGATCATTC AGCVATTGTG GGGAG 105

(2) INFORMATION FOR SEQ ID NO: 162:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 105 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 162:

CCCTTTAGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC GCCATACCGC AAGAGAGTCA 60
TGCCGCCTTT TTCACGACCG TGATCATTC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 163:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 163:

CCCTTTAGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGT GCCATACCGC AAGAGAGTCA	60
TGCCGCCTTT TTCACGACCG TGATCATTCC AGCCATTGTT GGGGG	105

(2) INFORMATION FOR SEQ ID NO: 164:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 164:

CCCTTTAGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGT GCCATACCGC AAGAGAGTCA	60
TGCCGCCTTT TTCACGACCG TGATCATTCC AGCCATTGTT GGGGG	105

(2) INFORMATION FOR SEQ ID NO: 165:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 165:

CCCTTTAGTT TCTCTCGTTT TAGCAGGAGC GTTGGTTAGT GCCATACCGC AAGAGAGTCA 60

TGCCGCCTTT TTCACGACCG TGATCATTCC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 166:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 105 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 166:

CCCTTTAGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGT GCCATACCGC AAGAGAGTCA 60

TGCCGCCTTT TTCACAACCG TGATCATTCC ARCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 167:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 105 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 167:

CCCTTTAGTT TCTCTTGTTT TAGCAGGAGC GTTGATTAGC GCCATACCGC AAGAGAGTCA 60

TGCCGCCTTT TTCACAACCG TAATCATTCC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 168:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 168:

CCCTTTAGTT TCTCTGTTT TAGCAGGAGC GTTGATTAGC GCCATACCGC AAGAGAGTCA 60

TGCCGCCTTT TTCACGACCG TAATCATTCG AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 169:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 169:

CCCTTTAGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC GCCATACCGC AAGAGAGTCA 60

TGCCGCCTTT TTCACAACCG TNATCATTCG AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 170:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 170:

CCCTTTGGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC TCCATACCGC AAGAGAGTCA 60
TGCCGCCTTT TTCACACCCG TGATCATTCC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 171:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 105 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 171:

CCCTTTGGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC TCCATACCGC AAGAGAGTCA 60
TGCCGCCTTT TTCACAACCG TGATCATTCC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 172:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 105 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 172:

CCCTTTGGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC TCCATACCGC AAGAGAGTCA 60
TGCCGCCTTT TTCACAACCG TGATCATTCC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 173:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 173:

CCCTTTGGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC TCCATACCGC AAGAGAGTCA	60
TGCCGCCTTT TTCACAACCG TGATCATTCC AGCCATTGTT GGGGG	105

(2) INFORMATION FOR SEQ ID NO: 174:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 174:

CCCTTTAGTT TCTCTCGTTT TAGCAGGAGC GTTGATTAGC TCCATACCGC AAGAGAGTCA	60
TGCCGCCTTT TTCACAACCG TGATCATTCC AGCCATTGTT GGGGG	105

(2) INFORMATION FOR SEQ ID NO: 175:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 175:

CCCTTTAGTT TCTCTCGTTT TAGCANNAGC GTTGATTAGC RCCATACCGC AAGAGAGTCA 60

TGCCGCCTTT TTTACAACCG TGATCATTCC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 176:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 105 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 176:

CCCTTTAGTT TCTCTTGTTT TAGCAGGAGC GTTGATTAGC GCCATACCGC AACAAAGTCA 60

TGCCGCCTTT TTCACGACCG TNATCATTCC AGCCATTGTT NNNNG 105

(2) INFORMATION FOR SEQ ID NO: 177:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 105 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 177:

CCCTCTGGTT TCTCTGGTTT TAGCAGGAGC GTTGATTAGC ATCACACCAC AACAAAGTCA 60

TGCCGCCTTT TTCACGACCG TGATCATTCC AGCCATTGTT GGGGG

105

(2) INFORMATION FOR SEQ ID NO: 178:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 178:

CCCTCTGGTT TCTCTCGCTT TAGTAGGAGC ATTGGTCAGC ATCACACCGC AACAAAGTCA 60

TGCCGCCTTT TTCACAACCG TGATCATTCC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 179:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 179:

CCCTCTGGTT TCTCTCGCTT TAGTAGGAGC ATTGGTCAGC ATCACACCGC AACAAAGTCA 60

TGCCGCCTTT TTCACAACCG TGATCATTCC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 180:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 180:

CCCTCTGGTT TCTCTCGCTT TAGTAGGAGC ATTGGTCAGC ATCACACCGC AACAAAGTCA 60

TGCCGCCTTT TTCACAACCG TGATCATTC AGCCATTGTT GGAGG 105

(2) INFORMATION FOR SEQ ID NO: 181:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 105 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 181:

CCCTCTGGTT TCTCTCGCTT TAGTAGGAGC ATTGGTCAGC ATCACACCGC AACAAAGTCA 60

TGCCGCCTTT TTTACAACCG TGATCATTC AGCCATTGTT GGAGG 105

(2) INFORMATION FOR SEQ ID NO: 182:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 105 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 182:

CCCTTTGGTT TCTCTCGCTT TAGTAGGAGC ATTGGTCAGC ATCACACCGC AACAAAGTCA 60

TGCCGCCTTT TTCACGACCG TGATCATTCC AGCCATTGTT GGGGG

105

(2) INFORMATION FOR SEQ ID NO: 183:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 183:

CCCTCTGGTT TCTCTTGCTT TAGTAGGAGC GTTAGTCAGC ATCACACCGC AACAAAGTCA 60

TGCCGCCTTT TTCACAACCG TGATCATTCC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 184:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 184:

CCCTCTAGTT TCTCTCGCTT TAGTAGGAGC ATTGGTCAGC ATCACACCGC AACAAAGTCA 60

TGCCGCCTTT TTCACAACCG TGATCATTCC AGCCATTGTG GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 185:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 185:

CCCTTTAGTT TCTCTTGCTT TAGTAGGAGC ATTGGTCAGC ATCACACCGC AACAAAGTCA 60

TGCCGCCTTT TTCACAACCG TGATCATTCC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 186:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 105 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 186:

CCCTCTNGTT TCTCTCGCTT TAGTAGNAGC ATTNGTCAGC ATCACACCGC AACANAGTCA 60

TGCCGCCTTT TTCACAACCG TNATCATTCC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 187:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 105 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 187:

CCCTCTGGTT TCTCTCGCTT TAGTAGGAGC GTTGGTCAGC ATCACACCGC AACAAAGTCA 60
 TGCCGCCTTT ATTACAACCG TGATCATTCC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 188:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 188:

CCCTCTGGTT TCTCTTGCTT TAGTAGGAGC ATTGGTCAGC ATCACACCGC AACAAAGTCA 60
 TGCCGCCTTT TTTACAACCG TGATTATTCC AGCCATTGTG GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 189:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 189:

CCCTCTGGTT TCTCTTGCTT TAGTAGGAGC ATTGGTCAGC ATCACACCGC AACAAAGTCA 60
 TGCCGCTCTT TTTACAACCG TGATTATTCC AGCCATTGTG GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 190:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 190:

CCCTCTGGTT TCTCTTGCTT TAGTAGGAGC ATTGGTCAGC ATCACACCGC AACAAAGTCA	60
TGCCGCTCTT TTTACAACCG TGATTATTCC AGCCATTGTG GGGGG	105

(2) INFORMATION FOR SEQ ID NO: 191:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 191:

CCCTCTGGTT TCTCTCGCTT TAGTAGGAGC ATTGGTCAGC ATCACACCGC AACAAAGTCA	60
TGCCGCTCTT TTTACAACCG TGATTATTCC AGCCATTGTG GGGGG	105

(2) INFORMATION FOR SEQ ID NO: 192:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 192:

CCCTYTGGTT TCTCTTGCTT TAGTAGGAGC ATTGRRTAGY RYCAYACCGC AACAAAGTCA 60
 TGCCGCCTTT TTYACRACCG TGATCATTC AGCCATTGTT GGRGG 105

(2) INFORMATION FOR SEQ ID NO: 193:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 105 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 193:

CCCTATTATC TCTCTCGCTT TAGTAGGAGC ATTGGTCAGC ATCACACCGC AACAAAGTCA 60
 TGCCGCCTTT TTCACAACCG TGTTTCATTCC AGCCATTGTT GGGGG 105

(2) INFORMATION FOR SEQ ID NO: 194:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 194:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA 60
 AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCACAAAGAG CGATAACGGG 120
 CTAAACACTA GCACTTTGGA TTTTAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC 180
 ACTACATCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT AGTGGTTACG 240
 ACCCGTGTTC AGAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT 300

CGCATTGGTG TCGTTAGTTT GCAAACGGG TATAGCCCGG CCTATTCTGG GGGCGTTACT 360

TT 362

(2) INFORMATION FOR SEQ ID NO: 195:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 195:

GTGGATGCTC ATACAGCTTA TTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA 60

AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCACAAAGAG CGATAACGGG 120

CTAAACACTA GTGCTTTGGA TTTGAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC 180

ACTACATCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT AGTGTTTACA 240

ACCCGTGTTC AGAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCG 300

CACATTGGTG TCGTTAGTTT GCAAACGGG TATAGCCCGG CCTATTCTGG GGGCGTTACT 360

TT 362

(2) INFORMATION FOR SEQ ID NO: 196:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 196:

GTGGATGCTC ATACAGCTTA TTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA 60

AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCACAAAGAG CGATAACGGG	120
CTAAACACTA GCGCTTTGGA TTTGAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACGTTAAA AACTTTGACA TTAAGGAATT GGTGGTTACA	240
ACCCGAGTTC AAAGTTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCG	300
CGCATTGGTG TCGTTAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 197:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 362 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 197:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCACAAAGAG CGATAACGGG	120
CTAAACACTA GCGCTTTGGA TTTGAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACGTTAAA AACTTTGACA TTAAGGAATT GGTGGTTACA	240
ACCCGAGTTC AAAGTTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCG	300
CGCATTGGTG TCGTTAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 198:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 362 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 198:

GTGGATGCTC ATACAGCTTA TTTTAATGGT AATATTTATC TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCACAAAGAG CGATAACGGG	120
CTAAACACTA GCGCTTTGGA TTTCAGCGGC GTTACAGATA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAACGT GAACATTAAA AACTTTGACA TTAAGGAATT GGTGGTTACA	240
ACCCGAGTTC AAAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTGAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 199:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 362 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 199:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGTAAGAG CGATAACGGG	120
CTAAACACTA GCACTTTGGA TTTTAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACGTTAAA AACTTTGACA TTAAGGAATT GGTGGTTACA	240
ACCCGAGTTC AAAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATCAGTCT	300
CGCATTGGTG TCGTGAGTTT GCAAACGGGA TATAGCCCGG CTTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 200:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 362 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 200:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CTAGTAAGAG CGATAACGGG	120
CTAAACACTA GCGCTTTGGA TTTTAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT GGTGGTTACA	240
ACCCGAGTTC AAAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATGGTG TCGTGAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 201:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 2 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 201:

WC

2

(2) INFORMATION FOR SEQ ID NO: 202:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 362 base pairs
 - (B) TYPE: nucleic acid

(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 202:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATGTTTATC TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGCAAGAG CGATAACGGG	120
CTAAACACTA GTGCTTTGGA TTTTAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT AGTGGTTACA	240
ACCCGAGTTC AAAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTGAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 203:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 362 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 203:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATGTTTATC TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGCAAGAG CGATAACGGG	120
CTAAACACTA GTGCTTTGGA TTTTAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT AGTGGTTACA	240
ACCCGAGTTC AAAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300

CGCATTGGTG TCGTGAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT 360
 TT 362

(2) INFORMATION FOR SEQ ID NO: 204:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 204:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATGTTTATC TGGGAAAATC CACGAATTTA 60
 AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGCAAGAG CGATAACGGG 120
 CTAAACACTA GTGCTTTGGA TTTTAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC 180
 ACTACATCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT AGTGGTTACA 240
 ACCCGAGTTC AAAGTTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT 300
 CGCATTGGTG TCGTGAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT 360
 TT 362

(2) INFORMATION FOR SEQ ID NO: 205:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 205:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATGTTTATC TGGGAAAATC CACGAATTTA 60

AGAGTGAATG CCCATAGCGC TCATTTTAAA AATATTGATG CCAGCAAGAG CGATAACGGG	120
CTAAACACTA GTGCTTTGGA TTTTAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT AGTGGTTACA	240
ACCCGAGTTC AAAGTTTTGG GCAATACACT ATTTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTGAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 206:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 362 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 206:

GTGGATGCTC ATACAGCTTA TTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCACAAAGAG CGATAACGGG	120
CTAAACACTA GCGCTTTGGA TTTTAGTGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT AGTGGTTACA	240
ACCCGAGTTC AAAGTTTTGG GCAATACACT ATTTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTTAGTTT GCAAACGGGA TATAGCCCAG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 207:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 362 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 207:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGTAAGAG CGATAACGGG	120
CTAAACACTA GTGCTTTGGA TTTTAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT AGTTGTTACA	240
ACCCGAGTTC AAAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTTAGTTT GCAAACGGGA TATAGCCCAG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 208:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 362 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 208:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AAAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGTAAGAG CGATAATGGT	120
CTAAACACTA GCGCTTTGGA TTTTAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT AGTGGTTACA	240
ACCCGTGTTC AGAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTTAGTTT GCAAATGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 209:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 362 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 209:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AAAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGTAAGAG CGATAACGGG	120
CTAAACACTA GCGCTTTGGA TTTGAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACAGCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT GGTGTTACG	240
ACCCGTGTTC AGAGTTTGG ACAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTTAGTTT GCAAACGGGA TATAGCCCGG CTTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 210:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 362 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 210:

GTGGATGCTC ATACAGCTAA CTTTAATGGC AATATTTATT TGGGAAAATC CACGAATTTA	60
AGGGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGTAAGAG CGATAACGGG	120
CTAAACACTA GCTCTTTGGA TTTCAGTGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACGTTAAA AACTTTGACA TTAAGGAATT GGTGGTTACA	240

ACCCGAGTTC AGAGTTTTGG GCAATACACT ATTTTTGGCG AAATTATAGG CGATAAGTCT	300
CGCATTGGTG TCGTTAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 211:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 211:

GTGGATGCTC ATACAGCTTA TTTAATGGC AATATTTATT TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGTAAGAG CGATAACGGG	120
CTAAACACTA GTGCTTTGGA TTTTAGCGGC GTTACAGATA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACGTTAAA AACTTTGACA TTAAGGAATT GGTGGTTACA	240
ACCCGAGTTC AAAGTTTTGG GCAATACACT ATTTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTTAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 212:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 212:

GTGGATGCCC ATACGGCTAA CTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AAAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCACAAAGAG CGATAACGGG	120
CTAAACACTA GCGCTTTGGA TTTGAGTGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT AGTGGTTACA	240
ACCCGTGTTC AGAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTTAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 213:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 213:

GTGGATGCTC ATACAGCTTA TTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AAAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGTAAGAG CGATAATGGT	120
CTAAACACTA GTGCTTTGGA TTTGAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACAGCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT AGTGGTTACG	240
ACCCGTGTTC AGAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG AGATCAATCG	300
CGCATTGGTG TCGTTAGTTT GCAAACGGG TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 214:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 214:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AAAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCACTAAGAG CGATAATGGT	120
CTAAACACTA GCGCTTTGGA TTTGAGCGGC GTTACAAACA AGGTCAATAT CAACAAGCTC	180
ACTACAGCTG CCACTAATGT GTCCATTAAA AACTTTGACA TTAAGGAATT AGTGGTTACG	240
ACCCGTGTTC AGAGTTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATCAATCG	300
CGCATTGGTG TCGTTAGTTT GCAAACGGC TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 215:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 362 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 215:

GTGGATGCTC ATACGGCTAA CTTTAATGGC AATGTTTATC TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGTAAGAG CGATAACGGG	120
CTAAACACTA GCGCTTTGGA TTTTAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT GGTGGTTACA	240
ACCCGTGTTC AGAGTTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTTAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 216:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 216:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATGTTTATC TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGCAAGAG CGATAACGGG	120
CTAAACACTA GCGCTTTGGA TTTTAGCGGC GTTACAGATA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT AGTGGTTACA	240
ACCCGTGTTC AGAGTTTTGG GCAATACACT ATTTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTGAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 217:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 217:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATGTTTATC TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CTAGTAAGAG CGATAACGGG	120

CTAAACACTA GCGCTTTGGA TTTTAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACGTTAAA AACTTTGACA TTAAGGAATT AGTGGTTACA	240
ACCCGTGTTT AGAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTGAGTTT GCAAACGGGA TATAGCCCAG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 218:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 362 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 218:

GTGGACGCTC ATACAGCTTA TTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGCAAGAG CGATAACGGG	120
CTAAACACTA GCGCTTTGGA TTTAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACGTTAAA AACTTTGACG TTAAGGAATT GGTGGTTACA	240
ACCCGTGTTT AGAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTGAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 219:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 362 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 219:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGCAAGAG CGATAACGGG	120
CTAAACACTA GCACTTTGGA TTTTAGCGGC GTTACAGATA AAGTCAATAT CAACAAGCTC	180
ACTACAGCTG CCACTAATGT GAACGTTAAA AACTTTGACA TTAAGGAATT AGTGGTTACA	240
ACCCGAGTTC AAAGTTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTTAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 220:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 220:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATT TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGTAAGAG CGATAACGGG	120
CTAAACACTA GCGCTTTGGA TTTTAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACGTTAAA AACTTTGACA TTAAGGAATT AGTGGTTACA	240
ACCCGTGTTT AGAGTTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTTAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 221:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid

(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 221:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATT TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGTAAGAG CGATAACGGG	120
CTAAACACTA GTGCTTTGGA TTTTAGCGGC GTTACAGATA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACGTTAAA AACTTTGACA TTAAGGAATT GGTGGTTACA	240
ACCCGAGTTC AAAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTTAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 222:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 362 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 222:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AGATTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGTAAGAG CGATAACGGG	120
CTAAACACTA GCGCTTTGGA TTTTAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACGTTAAA AACTTTGACA TTAAGGAATT GGTGGTTACA	240
ACCCGAGTTC AGAGTTTGG GCAATACTCT ATTTTGGCG AAAATATAGG CGATAAGTCG	300

CGCATTGGTG TCGTTAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT 360

TT 362

(2) INFORMATION FOR SEQ ID NO: 223:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 223:

GTGGACGCTC ATACAGCTTA TTTTAATGGC AATATTTATT TGGGAAAATC CACGAATTTA 60

AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCACAAAGAG CGATAACGGG 120

CTAAACACTA GCACTTTGGA TTTGAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC 180

ACTACATCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT AGTGGTTACG 240

ACCCGTGTTC AGAGTTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT 300

CGCATTGGTG TCGTGAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT 360

TT 362

(2) INFORMATION FOR SEQ ID NO: 224:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 224:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA 60

AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCACAAAGAG CGATAACGGG	120
CTAAACATTA GCACTTTGGA TTTTAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT AGTGGTTACA	240
ACCCGTGTTC AGAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTTAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 225:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 225:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTG	60
AGAGTGAATG GCCATAACGC TCATTTTAAA AATATTGATG CCAGTAAGAG CGATAACGGG	120
CTAAACACTA GCACTTTGGA TTTGAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACAGCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT AGTGGTTACG	240
ACCCGTGTTC AGAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG TGATAAGTCT	300
CGCATTGGTG TCGTTAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 226:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 226:

GTGGATGCTC ATACAGCTTA TTTAATGGC AATGTTTATT TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGTGAGAG CGATAACGGG	120
CTAAACACTA GCGCTTTGGA TTTTAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACGTTAAA AACTTTGACA TTAAGGAATT GGTGGTTACG	240
ACCCGTGTTT AGAGTTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTTAGTTT GCAAACGGGA TGTCGCCCCG CCTGTTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 227:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 227:

GGGGATGCTC ATACAGCTTA TTTAATGGC AATATTTTATT TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGTAAGAG CGATAACGGG	120
CTAAACACTA GCGYTTTGA TTTTAGCGGC GTTACAGAYA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACRTTAAA AACTTTGAYA TTAAGGAATT GGTGGTTACA	240
ACCCGAGTTC AAAGTTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATMAGTCT	300
CGCATTGGTG TCGTTAGTTT GCAAACGGGA TATAGCCCRG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 228:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 362 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 228:

GTGGATGCTC ATACAGCTTA TTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCACAAAGAG CGATAATGGT	120
ATAAACACTA GCACTTTGGA TTTGAGCGGC GTTACAGACA AGGTCAATAT CAACAAGCTC	180
ATTACAGCTT CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT GGTGGTTACA	240
ACCCGTGTTT AAAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATGGTG TCGTTAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 229:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 362 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 229:

GTGGATGCTC ATACGGCTAA CTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAA AATATTGATG CCAGCAAGAG CGATAACGGG	120
CTAAACACTA GCACCTTGA TTTAGTGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACGTTAAA AACTTTGATA TTAAGGAATT GGTGGTTACA	240

ACCCGAGTTC AGAGTTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCG	300
CGCATTGGTG TCGTGAGTTT GCAAACGGGA TATAGCCCAG CTTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 230:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 362 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 230:

GTGGATGCTC ATACGGCTAA CTTAATGGC AATATTTATT TGGGAAAATC CACGAATTTG	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGTAAGAG CGATAACGGG	120
CTAAACACTA GCGCTTTGGA TTTTAGCGGC GTTACAGACA AAGTTAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACGTTAAA AACTTTGACA TTAAGGAATT GGTGGTTACA	240
ACCCGAGTTC AAAGTTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCT	300
CGCATTGGTG TCGTGAGTTT GCAAACGGGA TATAGCCCTG CTTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 231:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 362 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 231:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AGAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCAGTAAGAG CGATAACGGG	120
CTAAACACTA CCACTTTGGA TTTCAGCGGC GTTACAGATA AAGTCAATAT CAACAAGCTC	180
ACTACATCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT AGTGGTTACA	240
ACCCGAGTTC AGAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGCTG	300
CACATTGGTG TCGTGAGTTT GCAAACGGGA TATAGCCCAG CCTATTCTGG GGGGCTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 232:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 232:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AAAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CTAGTAAGAG CGATAACGGG	120
CTAAACACTA GCGCTTTGGA TTGAGCGGC GTTACAAACA AGGTCAATAT CAACAAGCTC	180
ACTACAGCTG CCACTAATGT GAACATTAAA AACTTTGACA TTAAGGAATT GGTGGTTACA	240
ACCCGCGTTC AGAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATAAGTCG	300
CGCATTGGTG TCGTTAGTTT GCAAACGGG TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 233:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 233:

ATGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AAAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCACTAAGAG CGATAATGGT	120
CTAAACACTA GCGCTTTGGA TTTGAGCGGC GTTACAAACA AGGTCAATAT CAACAAGCTC	180
ACTACAGCTG CCACTAATGT GTCCATTAAA AACTTTGACA TTAAGGAATT AGTGGTTACG	240
ACCCGTGTTC AGAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATCAATCG	300
CGCATTGGTG TCGTTAGTTT GCAAACGGC TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 234:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 362 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 234:

GTGGATGCTC ATACAGCTTA TTTTAATGGC AATATTTATC TGGGAAAATC CACGAATTTA	60
AAAGTGAATG GCCATAGCGC TCATTTTAAA AATATTGATG CCACTAAGAG CGATAATGGT	120
CTATACACTA GCGCTTTGGA TTTGAGCGGC GTTACAAACA AGGTCAATAT TAACACGCTC	180
ACTACAGCTG CCACTAATGT GTCCATTAAA AACTTTGACA TTAAGGAATT AGTGGTTACG	240
ACCCGTGTTC AGAGTTTGG GCAATACACT ATTTTGGCG AAAATATAGG CGATCAATCG	300
CGCATTGGTG TCGTTAGTTT GCAAACGGC TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 235:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 235:

GTGGATGCCC ATACGATCAA TTTTAATGGC AATATGTATT TGGGAAGATT TACGCATTTA	60
AAAGTGAATG GTCATACAGC CAATTTTAAA GATATTGATG CCAGCAAGGG TAGAAATGGT	120
ATCGACACCA CCATTTTGGG TTTTAGCGGC GTTACAAACA AGGTCAATAT CAACAAGCTC	180
ACCACAGCTG CCACTAATGC GGCCATTAAA AATTTTGACA TTAAGGAATT GGTTGTTACA	240
ACCAATGTTT TGAGTGTGGG GAAATACACT GATTTTACCG AAGATATAGG CGATCAATCC	300
CGCATTGGTA TCGTGCCTTT GCAAATGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 236:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 236:

GTGGATGCCC ATACGATCAA TTTTAATGGC AATATGTATT TGGGAAGATT CACGCATTTA	60
AAAGTGAATG GTCATACAGC CAATTTTAAA GATATTGATG CCAGCAAGGG TAGAAATGGT	120

ATCGACACCA CCATTTTGGG TTTTAGCGGC GTTACAAACA AGGTCAATAT CAACAAGCTC	180
ACCACAGCTG CCACTAATGC GGCCATTAAA AATTTTGACA TTAAGGAATT GGTGTTACA	240
ACCAATGTTT TGAGTGTGGG GAAATACACT GATTTTACCG AAGATATAGG CGATCAATCC	300
CGCATTGGTA TCGTTAGTTT GCAAACGGGA TATAGCCCGG CCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 237:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 362 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 237:

GTGGATGCCC ATACGATCAA TTTTAATGGC AACATGTATT TGGGAAGATT CACGCATTTA	60
AAAGTGAATG GCCATACAGC CAATTTTAAA GATATTGATG CCAGCAAGGG TAGAAATGGT	120
ATCGACACCA CTATTTTGGG TTTTAGCGGC GTTACAGACA AAGTCAATAT CAACAAGCTC	180
ACTACAGCTG CCACTAATGT GTCCATTAAA AACTTTGACA TTAAGGAATT GGTGTTACA	240
ACCAATGTTT TGAGTGTGGG GAAATACACT GATTTTACCG AAGATATAGG CGATCAATCG	300
CACATTGGTG TCGTTAGTTT GCAAACGGG TATAGCCCGG TCTATTCTGG GGGCGTTACT	360
TT	362

(2) INFORMATION FOR SEQ ID NO: 238:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 238:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AGGTAAGGTC AATATCAACA AGCTCATTAC GGCTTCCACT	120
AATGTGGCCG CTAAAACTT CAACATTAAT GAATTGATTG TTTAAACCAA TGGGGTGAGT	180
GTGGGGGAAT AACTCATT TTAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG	240
CGTTTGGAAT CTGGCACTAG GTCAATCTAT TCTGGCGGTG TTAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 239:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 239:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AGGTAAGGTC AATATCAACA AGCTCATCAC AGCTTCCACT	120
AATGTGGCCG TTTAAAACTT CAACATTAAT GAATTGATTG TTTAAACCAA TGGTGTGAGT	180
GTGGGGGAAT AACTCATT TTAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG	240
CGTTTGGAAT CTGGCACTAG GTCAATCTTT TCTGGGGGTG TTAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 240:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 240:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AGGTAAGGTC AATATCAACA AGCTCATCAC AGCTTCCACT	120
AATGTGGCCG TTAAAACTT CAACATTAAT GAATTGATTG TTAAAACCAA TGGGATAAGT	180
GTGGGGGAAT ACACTCATTT TAGCGAAGAT ATAGGAAGTC AATCGCGCAT CAATACCGTG	240
CGTTTGAAA CTGGCACTAG ATCAATCTTT TCTGGGGGTG TTAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 241:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 241:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GCAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGCGTTAC AGACAAGGTC AATATCAACA AGCTCATTAC AGCTTCCACT	120
AATGTGGCCA TTAAAACTT CAACATTAAT GAATTGTTGG TTAAGACCAA TGGGGTGAGC	180
GTGGGGGAAT ACACTCATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG	240
CGTTTGAAA CTGGCACTAG GTCAATCTTT TCTGGGGGTG TCAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 242:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 242:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGCGTTAC AGACAAGGTC AATATCAACA AGCTCATTAC AGCTTCCACT	120
AATGTGGCCA TTAAAACTT CAACATTAAT GAATTGGTTG TTAAGACCAA TGGGGTGAGT	180
GTGGGGGAAT AACTCATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAACACCGTG	240
CGTTTGAAA CTGGCACTAG GTCAATCTAT TCTGGGGGTG TTAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 243:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 243:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AGGTAAGGTC AATATCAACA AGCTCATCAC GGCTTCCACT	120
AATGTGGCCG TTAAAAACAA CAACATTAAT GAATTGGTGG TTAAAACCAA TGGGATAAGT	180
GTGGGGGAAT AACTCATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG	240
CGTTTGAAA CAGGCACTAG GTCAATCTTT TCTGGGGGTG TCAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 244:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 244:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACTG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AAACAAAGTC AATATCAACA AGCTCATTAC AGCTTCCACT	120
AATGTGGCCG TTAAAACTT CAACATTAAT GAATTGTTGG TTAAGATTAA TGGGGTGAGT	180
GTGGGGGAAT ACACTTATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAACACCGTG	240
CGTTTGGAAG CTGGCACTAG GTCAATCTAT TCTGGCGGTG TTAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 245:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 245:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACTG GTAATGGTGG TTTCAACACC	60
TTAGATTTCA GTGGTGTTAC AGACAAGGTC AATATCAACA AGCTCATTAC AGCTTCCACT	120
AATGTGGCCA TTAAAACTT CAACATTAAT GAATTGGTGG TTAAAACCAA TGGTATAAGC	180
GTGGGGGAAT ACACTCATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAACACCGTG	240
CGTTTGGAAG CTGGCACTAG GTCAATCTAT TCTGGCGGTG TTAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 246:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 246:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACTG GTAATGGTGG TTTCAACACC	60
TTAGATTTCA GTGGTGTTAC AGACAAGGTC AATATCAACA AGCTCATTAC AGCTTCCACT	120
AATGTGGCCA TTAAAACTT CAACATTAAT GAATTGGTGG TTAAAACCAA TGGTATAAGC	180
GTGGGGGAAT ACACTCATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG	240
CGTTTGGAAG CTGGCACTAG GTCAATCTAT TCTGGCGGTG TTAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 247:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 288 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 247:

GTGGATGCCC ATACAGCTAA TTTTAAAGGT ATTGATACTG GTAATGGTGG TTTCAACACC	60
TTAGATTTCA GTGGCGTTAC AAACAAAGTC AATATCAACA AGCTCATTAC AGCTTCCACT	120
AATGTGGCCA TTAAAACTT CAACATTAAT GAATTGGTGG TTAAAACCAA TGGTATAAGC	180
GTGGGGGAAT AACTAATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAACACCGTG	240
CGTTTGGAAG CTGGCACTAG GTCAATCTAT TCTGGCGGTG TTAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 248:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 288 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 248:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AGGTAAGGTC AATATCAACA AGCTCATCAC AGCTTCCACT	120
AATGTGGCCG TTAAAACTT CAACATTAAT GAATTGATTG TTTAAACCAA TGGGGTGAGT	180
GTGGGGGAAT ACACTCATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG	240
CGTTTGAAAA CTGGCACTAG GTCAATCTAT TCTGGCGGTG TTAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 249:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 249:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AGGTAAGGTC AATATCAACA AGCTCATCAC AGCTTCCACT	120
AATGTGGCCG CTAAAACTT CAACATTAAT GAATTGATTG TTTAAACCAA TGGGGTGAGT	180
GTGGGGGAAT ACACTCATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG	240
CGTTTGAAAA CTGGCACTAG GTCAATCTAT TCTGGCGGTG TTAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 250:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 250:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AGGTAAGGTC AATATCAACA AGCTCATCAC AGCTTCCACT	120
AATGTGGCCG TTAAAACTT CAACATTAAT GAATTGATTG TTAAAACCAA TGGTGTGAGT	180
GTGGGGGAAT AACTCATT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAACACCGTG	240
CGTTTGAAA CTGGCACTAG GTCAATCTAT TCTGGCGGTG TTAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 251:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 288 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 251:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AGGTAAGGTC AATATTAACA AGCTCATTAC GGCTTCCACT	120
AATGTGGCCG TTAAAACTT CAACATTAAT GAATTGATTG TTAAAACCAA TGGGGTGAGT	180
GTGGGGGAAT AACTCATT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG	240
CGTTTGAAA CTGGCACTAG GTCAATCTTT TCTGGGGGTG TTAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 252:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 288 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 252:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTGAC AGGTATAGTC AATATCAACA AGCTCATCAC AGCTTCCACT	120
AATGTGGCCG TTAAAACTT CAACATTAAT GAATTGATTG TTAAAACCAA TGGGGTGAGC	180
GTGGGGGAAT ACACTCATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG	240
CGTTTGAAA CTGGCACTAG GTCAATCTTT TCTGGGGGTG TCAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 253:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 288 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 253:

GTGGATGGTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTAC AGGTAAGGTC AATATCAACA AGCTCATAAC GGCTTCCACT	120
AATGTGGCCG TTAAAACTT CAACATTAAT GAATTGATTG TTAAAACCAA TGGTGTGAGT	180
GTGGGGGAAT ACACTTATTT TAGCGAAGAT ATAGGCAGTC AATCGCACAT CAATACCGTG	240
CGTTTGAAA CTGGCACTAG GTCAATCTTT TCTGGGGGTG TCAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 254:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 288 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 254:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AGGTAAGGTC AATATCAACA AGCTCATTAC GGCTTCCACT	120
AATGTGGCCG TAAAAAACTT CAACATTAAT GAATTGATTG TTAAAACCAA TGGTGTGAGT	180
GTGGGGGAAT ACACTTATTT TAGCGAAGAT ATAGGCAGTC AATCGCACAT CAATACCGTG	240
CGTTTGAAA CTGGCACTAG GTCAATCTTT TCTGGGGGTG TCAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 255:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 288 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 255:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AGACAAAGTC AATATCAACA AGCTCATCAC AGCTTCCACT	120
AATGTGGCCG TAAAAAACTT CAACATTAAT GAATTGATTG TTAAAACCAA TGGGGTGAGT	180
GTGGGGGAAT ACACTCATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG	240
CGTTTGAAA CTGGCACCAG GTCAATCTTT TCTGGGGGTG TCAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 256:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 288 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 256:

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GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC      60
TTAGATTTTA GTGGTGTTAC AGACAAAGTC AATATCAACA AGCTCATTAC GGCTTCCACT      120
AATGTGGCCG TAAAAAAGTT CAACATTAAT GAATTGATTG TAAAAACCA TGGGGTGAGT      180
GTGGGGGAAT AACTCATT TTAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG      240
CGTTTGAAA CTGGCACTAG GTCAATCTTT TCTGGGGGTG TCAAATTT      288
```

(2) INFORMATION FOR SEQ ID NO: 257:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 288 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 257:

```
GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC      60
TTAGATTTTA GTGGTGTTAC AGACAAAGTC AATATCAACA AGCTCATTAC AGCTTCCACT      120
AATGTGGCCG TAAAAAAGTT CAACATTAAT GAATTGGTTG TAAAAACCA TGGGGTAAGT      180
GTGGGGGAAT AACTCATT TTAGCGAAGAT ATAGGCAGTC AATCGCACAT CAATACCGTG      240
CGTTTGAAA CTGGCACTAG GTCAATCTTT TCTGGGGGTG TCAAATTT      288
```

(2) INFORMATION FOR SEQ ID NO: 258:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 288 base pairs

(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 258:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AAACAAGGTC AATATCAACA AGCTCATTAC GGCTTCCACT	120
AATGTGGCCA TAAAAAATT CAACATTAAT GAATTGATTG TAAAACCAA TGGGATGAGC	180
GTGGGGGAAT AACTCATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG	240
CGTTTGAAA CTGGCACTAG GTCAATCTTT TCTGGGGGTG TCAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 259:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 288 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 259:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AAACAAGGTC AATATCAACA AGCTCATTAC AGCTTCCACT	120
AATGTGGCCG TAAAAAATT CAACATTAAT GAATTGATTG TAAAACCAA TGGGGTGAGC	180
GTGGGGGAAT AACTCATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG	240
CGTTTGAAA CTGGCACTAG GTCAATCTTT TCTGGGGGTG TCAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 260:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 288 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 260:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC 60
TTAGATTTTA GTGGTGTTAC AAACAAGGTC AATATCAACA AGCTCATTAC GGCTTCCACT 120
AATGTGGCCG TTAAAACTT CAACATTAAT GAATTGATTG TTAAAACCAA TGGGGTGAGC 180
GTGGGGGAAT AACTCATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG 240
CGTTTGAAA CTGGCACTAG GTCAATCTTT TCTGGGGGTG TCAAATTT 288

(2) INFORMATION FOR SEQ ID NO: 261:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 288 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 261:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC 60
TTGGATTTTA GTGGCGTTAC AGACAAAGTC AATATCAACA AGCTCATTAC AGCTTCCACT 120
AATGTGGCCA TTAAAACTT CAACATTAAT GAATTGTTGG TTAAGACCAA TGGGGTGAGT 180
GTGGGGGAAT AACTCATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAACACCGTG 240
CGTTTAGAAA CTGGCACTAG GTCAATCTTT TCTGGGGGTG TCAAATTT 288

(2) INFORMATION FOR SEQ ID NO: 262:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 288 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 262:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AGGTAAGGTC AATATCAACA AGCTCATTAC GGCTTCCACT	120
AATGTGGCCG TTAAAACTT CAACATTAAT GAATTGTTGG TTAAGACCAA TGGGGTGAGT	180
GTGGGGGAAT ACACTCATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG	240
CGTTTGAAAA CTGGCACTAG GTCAATCTTT TCTGGGGGTG TCAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 263:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 288 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 263:

GTGGATGCTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AGGTAAGGTC AATATCAACA AGCTCATTAC GGCTTCCACT	120
AATGTAGCCG TTAAAACTT CAACATTAAT GAATTGTTGG TTAAGACCAA TGGGGTGAGT	180
GTGGGGGAAT ACACTCATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAACACCGTG	240
CGTTTGAAAA CTGGCACTAG GTCAATCTTT TCTGGGGGTG TCAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 264:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 264:

GTGGATGGTC ATACAGCTAA TTTTAAAGGT ATTGATACGG GTAATGGTGG TTTCCACACC	60
TTAGATTTTA GTGGTGTTAC AGGTAAGGTC CATATCCACA AGCTCATTAC GGCTTCCACT	120
AATGTGGCCG TTAAAACTT CCACATTAAT GAATTGATTG GTAAAACCAA TGGGATAAGT	180
GTGGGGGAAT ACACTCATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG	240
CGTTTGGAAG CTGGCACTAG GTCAATCTTT TCTGGGGGTG TCAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 265:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 265:

GCGAGCGCTC ATACGGTCAA TTTTAAAGAT ATTGATACTG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AAACAAGGTC AATATCAACA AGCTCATTAC AGCTTCCACT	120
AATGTGGCCA TTAAAACTT CAACATTAAT GAATTGTTGG TTAAAACCAA TGGGATAAGT	180
GTGGGGGAAT AACTAATTT TAGCGAAGAT ATAGGCAATC AATCGCGCAT CAACACCGTG	240
CGTTTGGAAG CTGGCACCAG GTCAATCTAT TTTGGGGGTG TTAAATTA	288

(2) INFORMATION FOR SEQ ID NO: 266:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 266:

GCGAGCGCTC ATACGGTCAA TTTTAAAGAT ATTGATACTG GTAATGGTGG TTTCAACACC	60
TTAGACTTTA GTGGTGTTAC AAACAAGGTC AATATCAACA AGCTCATTAC AGCTTCCACT	120
AATGTGGCCG TTA AAAACTT CAACATTAAT GAATTGTTGG TTA AAACCA TGGGATAAGT	180
GTGGGGGAAT AACTAATTT TAGCGAAGAT ATAGGCAATC AATCGCGCAT CAACACCGTG	240
CGTTTGAAA CTGGCACTAG GTCAATCTAT TCTGGGGGTG TTAAATTT	288

(2) INFORMATION FOR SEQ ID NO: 267:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 267:

GCGAGCGCTC ATACGGTCAA TTTTAAAGAT ATTGATACTG GTAATGGTGG TTTCAACACC	60
TTAGACTTTA GTGGTGTTAC AAACAAGGTC AATATCAACA AGCTCATTAC AGCTTCCACT	120
AATGTGGCCA TTA AAAACTT CAACATTAAT GAATTGTTGG TTA AAACCA TGGGATAAGT	180
GTGGGGGAAT AACTAATTT TAGCGAAGAT ATAGGCAATC AATCGCACAT CAACACCGTG	240
CGTTTAGAAA CTGGCACTAG GTCAATCTAT TCTGGGGGTG TTAAGTTT	288

(2) INFORMATION FOR SEQ ID NO: 268:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 268:

GCGAGACGTC ATACGGTCAA TTTTAAAGAT ATTGATACTG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AAACAAGGTC AATATCAACA AGCTCATTAC AGCTTCCACT	120
AATGTGGCCG TTAAAACTT CAACATTAGT GAATTGTTGG TTAAAACCAA TGGGATAAGT	180
GTGGGGGAAT AACTAATTT TAGCGGAGAT ATAGGCAATC AATCGCGCAT CAACACCGTG	240
CGTTTGGAAT CTGGCACTAG GTCAATCTAT TCTGGGGGTG TTAAGTTT	288

(2) INFORMATION FOR SEQ ID NO: 269:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 269:

GTGGATGCCC ATACGGTCAA TTTTAAAGGT ATTGATACTG GTAATGGTGG TTTCAACACC	60
TTAGATTTCA GTGGTGTTAC AGACAAGGTC AATATCAACA AGCTCATTAC AGCTTCCACT	120
AATGTGGCCA TTAAAACTT CAACATTAAT GAATTGGTGG TTAAAACCAA TGGTATAAGC	180
GTGGGGGAAT AACTCATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG	240

CGTTTGGAAA CTGGCACTAG GTCAATCTAT TCTGGCGGTG TTAAATTT

288

(2) INFORMATION FOR SEQ ID NO: 270:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 270:

GCGAGCGCTC ATACGGTCAA TTTTAAAGAT ATTGATACTG GTAATGGTGG TTTCAACACC	60
TTAGACTTTA GTGGTGTTAC AAACAAGGTC AATATCAACA AACTCATTAC AGCTTCCACT	120
AATGTGGCCG TTAAAACTT CAACATTAAT GAATTGTTGG TTAAAACCAA TGGGATAAGT	180
GTGGGGGAAT AACTAATTT TAGCGAAGAT ATAGGCAATC AATCGCGCAT CAACACCGTG	240
CGTTTAGAAA CTGGCACCAG GTCAATCTAT TCTGGGGGTG TTAAGTTT	288

(2) INFORMATION FOR SEQ ID NO: 271:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 271:

GCGAGCGCTC ATACGGTCAA TTTTAAAGAT ATTGATACTG GTAATGGTGG TTTCAACACC	60
TTAGACTTTA GTGGTGTTAC AAACAAGGTC AATATCAACA AACTCATTAC AGCTTCCACT	120
AATGTGGCCG TTAAAACTT CAACATTAAT GAATTGTTGG TTAAAACCAA TGGGATAAGT	180
GTGGGGGAAT AACTAATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAATACCGTG	240

CGTTTAGAAA CTGGCACCAG GTCAATCTAT TCTGGGGGTG TTAAGTTT

288

(2) INFORMATION FOR SEQ ID NO: 272:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 272:

GCGAGCGCTC ATACGGTCAA TTTTAAAGAT ATTGATACTG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AAACAAGGTC AATATCAACA AGCTCATTAC AGCTTCCACT	120
AATGTGGCCA TTAAAACTT CAACATTAAT GAATTGTTGG TTAAAACCAA TGGGATAAGT	180
GTGGGGGAAT AACTAATTT TAGCGAAGAT ATAGGCAATC AATCGCGCAT CAATACCGTG	240
CGTTTAGAAA CTGGCACTAG GTCAATCTAT TCTGGGGGTG TTAAGTTT	288

(2) INFORMATION FOR SEQ ID NO: 273:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 273:

GCGAGCGCTC ATACGGTCAA TTTTAAAGAT ATTGATACTG GTAATGGTGG TTTCAACACC	60
TTAGATTTTA GTGGTGTTAC AAACAAGGTC AATATCAACA AGCTCATTAC AGCTTCCACT	120
AATGTGGCCG TTAAAACTT CAACATTAAT GAATTGTTGG TTAAAACCAA TGGGATAAGT	180

GTGGGGGAAT AACTAATTT TAGCGAAGAT ATAGGCAATC AATCGCGCAT CAACACCGTG	240
CGTTTGAAA CTGGCACTAG GTCAATCTAT TCTGGGGGTG TTAAGTTT	288

(2) INFORMATION FOR SEQ ID NO: 274:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 274:

GCGAGCGCTC ATACGGTCAA TTTTAAAGAT ATTGATACTG GTAATGGTGG TTTCAACACC	60
TTAGACTTTA GTGGTGTTAC AAACAAGGTC AATATCAACA AGCTCATTAC AGCTTCCACT	120
AATGTGGCCG TTAAAACTT CAACATTAAT GAATTGTTGG TTTAAACCAA TGGGATAAGT	180
GTGGGGGAAT AACTAATTT TAGCGAAGAT ATAGGCAATC AATCGCGCAT CAACACCGTG	240
CGTTTAGAAA CTGGCACTAG GTCAATCTAT TCTGGGGGTG TTAATTTT	288

(2) INFORMATION FOR SEQ ID NO: 275:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 288 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 275:

GCGAGCGCTC ATACGGTCAA TTTTAAAGAT ATTGATACTG GTAATGGTGG TTTCAATACC	60
TTAGACTTTA GTGGTGTTAC AAACAAGGTC AATATCAACA AGCTCATTAC AGCTTCCACT	120
AATGTGGCCA TTAAAAATTT CAACATTAAT GAATTGTTGG TTTAAACCAA TGGGATAAGT	180

GTGGGGGAAT AACTAATTT TAGCGAAGAT ATAGGCAGTC AATCGCGCAT CAACACCGTG 240
 CGTTTAGAAA CTGGCACTAG GTCAATCTAT TCTGGGGGTG TTAAGTTT 288

(2) INFORMATION FOR SEQ ID NO: 276:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 288 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 276:

ACGAGCGCTC ATACGGTCAA TTTTAAAGAT ATTGATACTG GTAATGGTGG TTTCAACACC 60
 TTAGACTTTA GTGGTGTTAC AAACAAGGTC AACATCAACA AGCTCATTAC AGCTTCCACT 120
 AATGTGGCCA TTAAAACTT CAACATTAAT GAGTTGTTGG TAAAAACCAA TGGGATAAGT 180
 GTGGGGGAAT AACTAATTT TAGCGAAGAT ATAGGCAATC AATCGCGCAT CAACACCGTG 240
 CGTTTAGAAA CTGGCACTAG GTCAATCTAT TCTGGGGGTG TTAAGTTT 288

(2) INFORMATION FOR SEQ ID NO: 277:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 21 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 277:

ATGGAAATAC AACAAACACA C 21

(2) INFORMATION FOR SEQ ID NO: 278:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 19 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 278:

CTGCTTGAAT GCGCCAAAC

19

(2) INFORMATION FOR SEQ ID NO: 279:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 21 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 279:

CACAGCCACT TTCAATAACG A

21

(2) INFORMATION FOR SEQ ID NO: 280:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 280:

CGTCAAAATA ATTCCAAGGG

20